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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:

APPLICANT: Maggio, John E.

APPLICANT: Maggio, John E.

APPLICANT: Maggio, John E.

APPLICANT: Magneth, Patrick W.

TITLE OF INVENTION: Labelled -Amyloid Peptide and Methods
TITLE OF INVENTION: Labelled -Amyloid Patrimer's Disease
NUMBER OF SEGUENCES: 3

CORRESPONDENCE ADDRESS: 3

ADDRESSEE Schwegman, Lundberg & Woessner, P.A.

STREET: 3500 IDS Center
CITY: Minneapolis
COMPUTER: MADDRESSE SCHWESSES

COMPUTER: TAR PROPONDENCE ADDRESSES

COMPUTER: TAR PROPONDENCE ADDRESSES

COMPUTER: TAR PROPONDENCE ADDRESSES

COMPUTER: TAR PROPONDENCE ADDRESSES

COMPUTER: Date PROPONDENCE ADDRESSES

ATTORNEY/AGENT INFORMATION:

MAMDE: Muting, Ann M.

REGISTRATION NUMBER: 33,977

RELECTORMUNICATION INFORMATION:

MANDE: MARKER TRANSPONDENCE INMEMBER: 600.226-US-01

TELECTORMUNICATION INFORMATION:

TELECTORMUNICATION ADDRESSES

TELECTORMUNICATION ADDRESSES

TELECTORMUNICATION TARGET ADDRESSES

TELETRANCE ADDRE
                             Sequence
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                                               US-08-404-831-1
US-08-60-2244A-3
US-08-62-362-1
US-08-61-765B-1
US-08-91-731E-70
US-08-91-31E-70
US-08-91-31E-70
US-08-91-31E-70
US-08-91-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/07744767A; Patent No. 5434050; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPAX: 612.339-3061
| INFORMATION FOR SEQ ID NO. 2:
| SEQUENCE CHARACTERISTICS:
| LENGTH: 42 amino acids
| TYPE: amino acid
| TOPOLOGY: linear
| MODECULE TYPE: peptide
US-07-744-767A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
Matches 42; Conserv
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US-07-744-767A-2
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Sequence 1, Appli
Sequence 19, Appl
Sequence 19, Appl
Sequence 19, Appl
Sequence 19, Appl
Sequence 1, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 3, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 5, Appli
Sequence 6, Appli
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                                                                                                                                                                  April 21, 2003, 12:06:35; Search time 15 Seconds (without alignments)
82,384 Million cell updates/sec
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217
1 DAEFRHDSGYEVHHQKLVFF......DVGSNKGAIIGLMVGGVVIA 42
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3: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
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6: /cgn2_6/ptodata/1/iaa/pcTUS COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/pcTUS COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/pcTUS COMB.pep:*
                             GenCore version 5.1.4 p5 4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                           OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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Match Length
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APPLICATION NUMBER: US/08/347,144
                                                          ATTORNEY/GENT INFORMATION:
NAME: AUGREACH, UFFREY I.
REGISTRATION NUMBER: 32,680
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 383-7451
TELEFEX: (202) 383-7451
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 42 amino acids
TYPE: amino acids
TYPE: amino acids
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
CRIGINAL SOURCE:
ORGANISM: AMYLOID PEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-462-859A-19
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US~08-462-859A-19
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Sequence 1, Application US/08347144

Patent NO. 5599154

GENERAL INFORMATION:

TITLE OF INVENTION: METHODS FOR THE PREVENTION AND TREATMENT

TITLE OF INVENTION: OF VASCULAR HEMORRHAGING AND ALZHEIMER'S DISEASE

NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESSE:

ADDRESSEE: HOWER'& SIMON

STREET: 1299 PENNSYLVANIA AVENUE, N.W.

CITY: MASHINGTON

STATE: D.C.

COUNTRY: US

ZIP: 20004

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPAtible

COMPUTER: IBM PC COMPAtible

COMPUTER: IBM PC COMPATIBLE

COMPUTER: IBM PC COMPATIBLE

COMPUTER: PLOSS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25
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                                         USTURENT NAPPLICATION US/08179574

SEQUENCE 1/92-5/4.

PAPLICANT: MINICIPATION:

APPLICANT: USEMEN KAYYALI

TITLE OF INVENTION:

NUMBER OF SEQUENCES:

CORRESPONDENCE HAMILTON, Brook, Smith & Reynolds, P.C.

STREET: Two Militia Drive

CITY: Lexington

STREET: Two Militia Drive

CONTRY: Lexington

CONPUTER: Massachusetts

CONPUTER: Massachusetts

CONPUTER: Massachusetts

CONPUTER: Dexington

STREET: Two Militia Drive

CONPUTER: Ploppy disk

CONPUTER: Dexington

STREET: Application Drive

CONPUTER: ISM PC Comparible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CLASSIFICATION DATA:

APPLICATION NUMBER: US/08/179,574

FILING DATE:

CLASSIFICATION NUMBER: US/08/179,574

FILING DATE:

TELECOMMUNICATION NUMBER: H199.

ATTORNEY AGENT INFORMATION:

NAME: Granahan, Patricia

REGERERENCE/DOCKET NUMBER: H190-03A3

TELECOMMUNICATION NUMBER: H190-03A3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 217; DB 1; Length 42; Best Local Similarity 100.0%; Pred. No. 46-26; Matches 42; Conservative 0; Mismatches 0; Indels
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; INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
; ENGTR: 42 amino acids
; TYPE: amino acid
;; TOPOLOGY: linear
US-08-179-574-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ઠે
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us-09-580-018-42.rai

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APPLICANT: Jacobsen, J. S.
TITLE OF INVENTION: No. 5703209e1 Amyloid Precursor and Method of TITLE OF INVENTION: No. 5703209e1 Amyloid Peptide
TITLE OF INVENTION: Of B-Amyloid Peptide
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSER: American Cyanamid Company
STREET: One Cyanamid Plaza
CITY: MAYNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match

Query Match

100.0%; Score 217; DB 1; Length 42;

Best Local Similarity 100.0%; Preed, No. 4e-26;

Matches 42; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                    CURPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TISM FC Comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATORIEN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,247A
FILING DATE: 05-UN-1995
CLASSIFICATION 1435
ATTORNEY AGENT INPORMATION:
NAME: BATTMARC, Elizabeth M.
REGISTRATION NUMBER: 31,088
TELECOMMINICATION INDERR: 31,088
TELECOMMINICATION INPORMATION:
TELECHONE: 201-683-2158
TELECHONE: 201-683-4117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Wayne
CUNTRY: Wayne
COUNTRY: United States
ZIP: 07470-8426
ZIP: 07470-8426
COMPUTER: Bandable Form:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/464,248A
FILING DATE: 05-JUN-1995
CLASSIFICATION NUMBER: 31,098
ATTORNEY/AGENT INPORMATION:
REGISTRATION NUMBER: 31,098
REFERENCE/DOCKET NUMBER: 31,098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMYGGVVIA 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 201-683-4117
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 amino acids
            One Campus Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 42 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                  Parsippany
New Jersey
Y: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-464-247A-19
                                                                                                                      COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Д
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US-68-129

US-68-124

US-68-124

US-68-124

US-68-124

US-68-125

APPLICANT: Jacobsen, J. S.

APPLICANT: Jacobsen, J. S.

TITLE OF INVENTION: Using Same to Access Agents Which Down-Regulate Formation ITITLE OF INVENTION: Using Same to Access Agents Which Down-Regulate Formation ITITLE OF INVENTION: Using Same to Access Agents Which Down-Regulate Formation ITITLE OF INVENTION: Using Same to Access Agents Which Down-Regulate Formation ITITLE OF INVENTION: Using Same to Access Agents Which Down-Regulate Formation ITITLE OF INVENTION: Using Same to Access Agents Which Down-Regulate Formation ITITLE OF INVENTION: Using Same to Access Agents Which Down-Regulate Formation: Same to Comparible 
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Sequence 10. 569347B

Patent No. 569347B

GENERAL INNORMATION:

APPLICANT: Jacobsen, J. S.

APPLICANT: Vitek, M. P.

TITLE OF INVENTION: No. 5693478el Amyloid Precursor and Method of

TITLE OF INVENTION: Using Same to Access Agents Which Down-Regulate Formation

TITLE OF INVENTION: of B-Amyloid Peptide

NUMBER OF SEQUENCES: 19

CORRESPONDENCE ADDRESS:

ADDRESSEE: American Cyanamid Company
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Ouery Match
100.0%; Score 217; DB 1; Length 42;
Best Local Similarity 100.0%; Pred. No. 4e-26;
Matches 42; Conservative 0; Mismatches 0; Indels
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MOLECULE TYPE: protein
US-08-123-659A-19
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US-08-464-247A-19
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GENERAL INFORMATION:

APPLICANT: Madgio, John E.

APPLICANT: Mantyh, Patrick M.

TITLE OF INVENTION: METHODS FOR USE IN DETECTING ALZHEIMER'S DISEASE
TITLE OF INVENTION: METHODS FOR USE IN DETECTING ALZHEIMER'S DISEASE
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Methods Rasch, Gebhardt & Schwappach, P.A.
STREET: P.O. Box 581415
COUNTRY: Minneapolis
STRATE: MN
COUNTRY: Minneapolis
STRATE: MN
COUNTRY: USA
ZIPS Floppy disk
MEDIUM TYPES: Ploppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: 12-SEP-1994
CLASSIPICATION UNMBER: US/08/304,585
ATTORNEY/AGRATINN: MNABER: U10.00010120
TELECOMMUNICATION INPORMATION:
NAME: Museting, Ann M.
REGISTRATION NUMBER: 110.00010120
TELECOMMUNICATION INPORMATION:
TELEFAX: 612-305-1228
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 amino acide
TTELEFAX: G12-305-1228
TTELEFAX: G12-305-1228
TTELEFAX: C12-305-1228
TTELEFAX: C12-305-1238

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100.0%; Score 217; DB 1; Length 42;
Best Local Similarity 100.0%; Pred. No. 4e-26;
Matches 42; Conservative 0; Mismatches 0; Indels
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US-06-302-808-5

Sequence 5, Application US/08302808

Patent No. 576349

GENERAL INFORMATION:
APPLICANT: SUZUKI, No. 5750349uhiro
APPLICANT: SUZUKI, No. 5750349uhiro
APPLICANT: ODARA, Asano
APPLICANT: ODARA, Asano
APPLICANT: ODARA, Asano
APPLICANT: ODARA, Asano
APPLICANT: WANTIBODIES TO B-AMYLOIDS OR THEIR
TITLE OF INVENTION: DERIVATIVES AND USE THEREOF
NUMBES OF SUGUENCES: 14
CORRESPECT: 130 WATER STREET
ADDRESSED: DIKE, BRONSTEIN, ROBERTS & CUSHMAN
STREET: 130 WATER STREET
CITY: MA
COUNTRY: MA
COMPUTER: BASING COMPUTER: IBM COMPUTER: US/O8/302,808
FILING DATE: 15-SEP-1994
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not relevant
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MOLECULE TYPE: peptide

US-08-304-585-2
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Sequence 1, Application US/08476464A

Patent No. 5707821

GENERAL INFORMATION:

APPLICANT: RYDEL RUSSELL E.

APPLICANT: DAPPEN, MICHAEL S.

TITLE OF INVENTION: THERAPEUTIC INHIBITION OF PHOSPHOLIPASE

TITLE OF INVENTION: THERAPEUTIC INHIBITION OF PHOSPHOLIPASE

TITLE OF INVENTION: THERAPEUTIC INHIBITION OF PHOSPHOLIPASE

TITLE OF INVENTION: AZ IN A-BETA FEPTIDE-MEDIATED NEURODEGENERATIVE DISEASE

NUMBER OF SEQUENCES: 2

COMPRESSEDE TOWNSEND & TOWNSEND & CREW LLP

STREET: TWO EMBRESS:

COMPUTER: RAPACISCO

STATE: CALIFORNIA

COMPUTER: TWO EMBRESS:

COMPUTER: 18 PC COMpatible

OPERATING SYSTEM: PC COMpatible

OPERATING SYSTEM: PC DAMPS NAS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURSENT PAPLICATION NUMBER: US/08/476,464A

ATTORNEY/AGENT INFORMATION:

NAME: STORELLA, OCHN N.

RECISTRATION NUMBER: 12,344

ATTORNEY/AGENT INFORMATION:

NAME: STORELLA, OCHN N.

TELECOMMUNICATION NUMBER: 15,294

TELECOMMUNICATION PROPRIESTICS:

SEQUENCE REPRESENCE THE PROPRIESTICS:

SECUENCE REPRESENCE THE PROPRIESTICS:

SECUENCE REPRESENCE THE PROPRIESTICS:

SECUENCE REPRESE
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                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 217; DB 1; Length 42; Best Local Similarity 100.0%; Pred. No. 4e-26; Matches 42; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGALIGLMVGGVVIA 42
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                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 amino acida
TYPE: amino acid
STRANDEDNESS: ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 42 amino acids
TYPE: amino acid
                                                                                                                                                                                                                              TOPOLOGY: linear; MOLECULE TYPE: protein US-08-464-248A-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
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RESULT 9 US-08-304-585-2 , Sequence 2, Application US/08304585 , Patent No. 5721106

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Gaps

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US-08-433-734-2
; Sequence 2, Application US/08433734
; Patent No. 5837473
; GENERAL INFORMATION:
APPLICANT: Madgio, John E.
APPLICANT: Mantyi, Patrick W.
TITLE OF INVENTION: Labelled -Amyloid Peptide and Methods
SEQUENCE:
STREET: P.O. Box 581415
; CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
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100.0%; Score 217; DB 1; Length 42;
Best Local Similarity 100.0%; Pred. No. 4e-26;
Matches 42; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 4e-26;
Matches 42; Conservative 0; Mismatches 0; Indels
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STATE: MA

COUNTRY: USA

ZIIP: 55458-1415

COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PREADABLE
COMPUTER: PROPED MISK
COMPUTER: MAX-1995
CLASSIFICATION NUMBER: US/08/433,734
FILING DATE: US/08/43,734
FILING DATE: US/08/433,734
FILING DATE: US/08/403
FILING DATE: US/08/43,734
FILING DATE: US/08
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                     APPLICATION NUMBER: BP 93110755.1
FILING DATE: 06-JUL-1993
ATTORNEY/AGENT INFORMATION:
NAME: PATISE, JOHN P.
REGISTRATION NUMBER: 34-403
REFERENCE/DOCKET NUMBER: 4105/157
TELECOMMUNICATION INPORMATION:
TELEPHONE: (201) 235-5326
INFORMATION POR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 amino acid
TYPE: amino acid
TYPE: amino acid
TYPE: Alleant TYPE: Number Action acide
TYPE: Alleant TYPE: Peptide
FRAGMENT TYPE: Nuterminal
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URS-08-268-348A-1
Sequence I, Application US/08268348A
Fatent No. 578374
GENERAL INFORMATION:
FAPLICANT: Debeli, Heinz
APPLICANT: Trettman, Gerda H
APPLICANT: Trettman, Gerda H
APPLICANT: Stuber, Dietrich
TITLE OF INVENTION: Process for Producing Hydrophobic
TITLE OF INVENTION: Process for Producing Hydrophobic
TITLE OF INVENTION: Process for Producing Same
APPLICANT: Stuber, Dietrich
TITLE OF INVENTION: Producing Same
APPLICANT: Along Invention: Producing Same
TITLE OF INVENTION: Producing Same
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: ADOFTMESS:
ADDRESSEE: Hoffmann-La Roche Inc.
STREET: 340 Kingsland Street
CITY: Nutley disk
COMPITER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPITER: IBP PC compatible
COMPITER: Present IN PRESENT OF STREET:
APPLICATION NUMBER: US/08/268,348A
FILING DATE: 29-UDN-1994
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/JP94/00069
FILING DATE: 24-JAN-1993
FILING DATE: 25-JAN-1993
FILING DATE: 25-JAN-1993
FILING DATE: 05-FBB-1993
APPLICATION NUMBER: 019015/1993
FILING DATE: 16-NOV-1993
APPLICATION NUMBER: 286985/1993
FILING DATE: 16-NOV-1993
APPLICATION NUMBER: 34473/1993
FILING DATE: 28-DEC-1993
APPLICATION NUMBER: 344,235
FILING DATE: 28-DEC-1993
APPLICATION NUMBER: 34,235
FILING DATE: 28-DEC-1993
APPLICATION NUMBER: 34,235
FILING DATE: 28-DEC-1993
APPLICATION NUMBER: 34,235
FEGENENCE/DOCKET NUMBER: 44631
TELEFONE: 617-523-6440
TELEFONE: 8EQUENCE CHARACTERISTICS:
FUNDAMETRIFICS: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 42 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-terminal
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; ORIGINAL SOURCE:
US-08-302-808-5
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100.0%; Score 217; DB 2; Length 42;
Best Local Similarity 100.0%; Pred. No. 4e-26;
Matches 42; Conservative 0; Mismatches 0; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FESSEED for Windows Version 2.0
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/01/137,371E
FILING DATE: 29-UL-1991
CLASSIFICATION 536
PRIOR APPLICATION DATA:
APPLICATION UNMER: 07/559,172
FILING DATE: 27-UL-1990
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      00108/028002
                                                                                                                                                                                                                                                                                                                                       MAME: Freeman, John W.
REGISTRATION NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 0010
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-642-8906
TELERA: 2011544-8906
INFORMATION FOR SEQ ID NO: 72: SEQUENCE CHARACTERISTICS:
LENGTH: 42 amino acids
TYPE: amino acid TYPE: amino acids
TYPE: amino acid
TYPE: amino acid
TYPE: 1 inear
MOLECULE TYPE: protein
US-07-737-311E-72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ð
                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: HENSLEY, Kenneth
APPLICANT: HENSLEY, Kenneth
APPLICANT: HENSLEY, Kenneth
APPLICANT: BUTTERFIELD, D. A.
APPLICANT: GARNEY, John M.
APPLICANT: AKSENOV, Michael
ITILE OF INVENTION: AN OLIGOPEPTIDE OR POLYPEPTIDES
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: LOWE PRICE LEBLANC & BECKER
STREET: 99 Canal Center Plaza, Suite 300
CITY: Alexandria
COUNTRY: USA
ZIP: 22314
COMPUTER: FLOME PRICE LEBLANC & BECKER
STATE: Virginia
COUNTRY: USA
ZIP: POPPY disk
COMPUTER: FLOME PRICE LEBLANC BELAGE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Batentin Relasse #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/609,090
FILING DATE: 29-FEB-1996
CLASSIFICATION NUMBER: 36,190
REFERENCE/DOCKET NUMBER: 36,110
TELEFONE: 703-684-1111
TELEFONE: 703-684-1111
TELEFONE: AMINO ACIGGE
LENGTH: 42 amino acide
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100.0%; Score 217; DB 2; Length 42;
Best Local Similarity 100.0%; Pred. No. 4e-26;
Matches 42; Conservative 0; Mismatches 0; Indels
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Sequence 12, Application US/07737371E
Patent No. 5876948
GENERAL INFORMATION:
APPLICANT: Yankner, Bruce A.
TITLE OF INVENTION: SCREENING METHODS TO IDENTIFY
TITLE OF INVENTION: NEUROTOXIN INHIBITORS (AS AMENDED)
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESSE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
  1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , 4e-26;
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                                                                                                                                                          Sequence 9, Application US/08609090; Patent No. 5840838; GENERAL INFORMATION;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: peptide US-08-609-090-9
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US-07-737-371E-72
                                                                                                           RESULT 13
US-08-609-090-9
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WESULE 12-333-4
Sequence 4, Application US/08422333
Patent No. 5912410

APPLICANT: CORDELL, Barbara L.
TITLE OF INVENTION: THE ANTIOID-FORMING PATHOLOGY OF ALZHEIMER'S DISBASE NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: 2450 Bayehore Parkway
CITY: Mountain View
STREET: 2450 Bayehore Parkway
CITY: Mountain View
ASTREE: ADDRESS:
ADDRESSEE: Plopy disk
COUNTRY: USA
ZIP: 94043
ZIP: 94
Gaps
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COUNTRY: US ZIP: 02110-2804

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; STRANDEDNESS; single
; TOPOLOGY: linear
US-08-422-333-4
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Query Match 100.0%; Score 217; DB 2; Length 42; Best Local Similarity 100.0%; Pred. No. 4e-26; Matches 42; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Search completed: April 21, 2003, 12:08:07 Job time : 16 secs

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Sequence 1, Appli Sequence 1, Appli Sequence 5, Appli Sequence 173, Appl Sequence 173, Appl Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 10, Appl Sequence 10, Appl Sequence 10, Appl Sequence 11, Appl Sequence 11, Appl Sequence 13, Appl Sequence 13, Appl Sequence 13, Appl Sequence 14, Appl Sequence 16, Appl Sequence 17, Appl

Sequence 10, Sequence 12, Sequence 14, Sequence 14,

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0 US-09-996-357-1
0 US-09-994-894-1
0 US-09-994-894-1
0 US-09-97-543-5
0 US-09-155-076-14
0 US-09-155-076-14
0 US-09-155-076-14
0 US-09-125-076-14
0 US-09-125-10
0 US-09-126-10
0 US-09-104-975-10
0 US-09-104-977-10
0 US-09-104-977-10
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0 US-09-104-977-10
0 US-09-105-847-10
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   Sequence 2, Appli
Sequence 7, Appli
Sequence 37, Appl
Sequence 174, Appl
Sequence 174, Appl
Sequence 1, Appli
Sequence 3, Appli
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Sequence 3, Appli
Sequence 1, Appli
                                                                         April/21, 2003, 12:06:50 ; Search time 15 Seconds (without alignments) 211.719 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Description
                                                                                                                      US-09-580-018-42
217
1 DAEFRHDSGYEVHHQKLVFF.....DVGSNKGAIIGLMVGGVVIA
                                                                                                                                                                                                                                                                                                                       Published Applications AA:*
| cgn2 6/prodata/2/pubpaa/USO8 NEW PUB.pep:*
2: /cgn2 6/prodata/2/pubpaa/USO8 NEW PUB.pep:*
3: /cgn2 6/prodata/2/pubpaa/USO8 NEW PUB.pep:*
4: /cgn2 6/prodata/2/pubpaa/USO7 NEW PUB.pep:*
5: /cgn2 6/prodata/2/pubpaa/USO7 NEW PUB.pep:*
6: /cgn2 6/prodata/2/pubpaa/USO7 NEW PUB.pep:*
7: /cgn2 6/prodata/2/pubpaa/USO7 NEW PUB.pep:*
8: /cgn2 6/prodata/2/pubpaa/USO8 NEW PUB.pep:*
9: /cgn2 6/prodata/2/pubpaa/USO8 PUBCOMB.pep:*
10: /cgn2 6/prodata/2/pubpaa/USO8 PUBCOMB.pep:*
11: /cgn2 6/prodata/2/pubpaa/USO8 PUBCOMB.pep:*
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11: /cgn2 6/prodata/2/pubpaa/USO8 PUBCOMB.pep:*
13: /cgn2 6/prodata/2/pubpaa/USO8 PUBCOMB.pep:*
14: /cgn2 6/prodata/2/pubpaa/USO8 PUBCOMB.pep:*
            GenCore version 5.1.4 p5 4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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US-10-011-496-2

US-10-962-956-37

US-09-962-956-37

US-09-865-294-65

US-09-867-847-1

US-09-867-847-1

US-09-867-847-1

US-09-956-655-26

US-09-956-655-26

US-10-071-459-1

US-10-071-459-1

US-09-908-037-3

US-09-908-037-3
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                                                                                                                                                                                                    288829 seqs, 75613885 residues
                                                                                                                                                                                                                                                                            Post-processing: Minimum Match 10%
Maximum Match 100%
Listing first 45 summaries
                                                       OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Perfect score:
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Publication No. US20030013648A1
GENERAL INFORMATION:
APPLICANT: Gerardo M. Castillo
APPLICANT: Gerardo M. Castillo
MUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSES: Patrick M. Dwyer
STREET: ProteoTech, Inc, 1818 Westlake Avenue N, Suite 114
CITY: Seattle
STARET: WA (Washington)
COUNTRY: United States of America
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COMPUTER READBLE FORM:
COMPUTER: 1BM PC
COMPUTER: 1BM PC
COMPUTER: 1 BM PC
COMP
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APPLICANT: Gurney, Mark E.
ITILE OF INVENTION: Transgenic Knockouts of BACE-1
FILE REFERENCE: MBHB 02-329-8
CURRENT APPLICATION WINBER: US/10/082,804
CURRENT APPLICATION WINBER: 60/271,092
PRIOR PLING DATE: 2001-02-23
PRIOR PLING DATE: 2001-02-26
PRIOR PLING DATE: 2001-02-26
PRIOR PLING DATE: 2001-02-26
PRIOR PLING DATE: 2001-05-26
PRIOR PLING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.1
                                                                                                                                        1 DAEFRHDSGYEVHHQKLVFFABDVGSNKGAIIGLMVGGVVIA 42
                                                                                                      1 DAEFRHDSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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   Best Local Similarity 100.0%; Pred. No. 1.5e-22; Matches 42; Conservative 0; Mismatches 0;
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100.0%; Score 217; DB 9;
Best Local Similarity 100.0%; Pred. No. 1.5e-22;
Matches 42; Conservative 0; Mismatches 0;
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; OTHER INFORMATION: A-beta 42 sequence.
US-10-082-804-7
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CLASSIFICATION
PRIOR APPLICATION DATE:
APPLICATION NUMBER: 09/938,275
FILING DATE: 22-August-2001
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                          ; Sequence 7, Application US/10082804; Publication No. US20020194632A1; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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US-09-962-955C-37
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US-10-082-804-7
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Publication No. US20020182660A1
GENERAL INFORMATION:
APPLICANT: Kei-Lai L. Fong
TITLE OF INVENTION: N- and C-Terminus Specific Immunoassays for
Abeta(1-39)
Abeta(1-41), Abeta(1-42) and Abeta (1-43)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity to other sequences, hydro-phobic
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                                                                    Length 42;
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                                                                                                                                                                                              1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGALIGLMVGGVVIA 42
                                                                                                                                                                                                                                               1 DAEFRHOSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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MEDIUM TYPE: 3.50 inch, 1.44MB storage COMPUTER: 1BM PC Compatibles
COMPUTER: 1BM PC Compatibles
OPERATING SYSTEM: Windows
SOFTWARE: MS No. US20020182660Alepad
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION TOWNER: US/10/051,496
FILING DATE: 18.50m-2002
CLASSIFICATION: UNKNOWN-
PRIOR APPLICATION TOWNER: US/09/784,854A
FILING DATE: 16-Feb-2001
APPLICATION NUMBER: 05/183,407
FILING DATE: 18-February-2000
ATTORNEY/AGENT INFORMATION:
                                                            Query Match
100.0%; Score 217; DB 8;
Best Local Similarity 100.0%; Pred. No. 1.5e-22;
Matches 42; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Koenig, C. Frederick III
REGISTRATION NUMBER: 29,662
REFERENCE/DOCKET NUMBER: PBI-PT001.1
TELECOMMUNICATION INFORMATION:
TELEFANK: (215) 568-6499
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSER: Kei-Lai L. Fong
STREET: 1004 West &th Avenue
CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Signal Sequence
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IDENTIFICATION METHOD:
OTHER INPORMATION:
AUTHORS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
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US-10-051-496-2
US-08-923-055-2
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JOTHER INFORMATION: Description of Artificial Sequence: All D peptides

JOTHER INFORMATION: or peptidomimetics

JOS-09-867-847-1
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100.0%; Score 217; DB 10; Length 42;
Best Local Similarity 100.0%; Pred. No. 1.5e-22;
Matches 42; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                            Length 42;
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100.0%; Score 217; DB 9; Sest Local Similarity 100.0%; Pred. No. 1.5e-22; Matches 42; Conservative 0; Mismatches 0;
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ORGANISM: Artificial Sequence
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CRGANISM: Homo sapiens

US-09-956-625-26
     ; SEQ ID NO 65
; LENGTH: 42
; TYPE: PRT
; ORGANIEM: Homo sapiens
US-09-865-294-65
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US-09-956-625-26
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Publication No. US20030068325A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Wang Yi
TITLE OF INVENTION: Immunogenic peptide composition as vaccines for the
TITLE OF INVENTION: Immunogenic peptide composition as vaccines for the
TITLE OF INVENTION: prevention and treatment of Alzheimer's Disease
FILE REFERENCE: 1151-4167
CURRENT APPLICATION NUMBER: US/09/865,294
CURRENT APPLICATION NUMBER: 2010-05-25
NUMBER OF SEQ ID NOS: 76
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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100.0%; Score 217; DB 9;
Best Local Similarity 100.0%; Pred. No. 1.5e-22;
Matches 42; Conservative 0; Mismatches 0;
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) OTHER INFORMATION: Amyloid Beta Peptide

US-09-848-616-174
NAME: DWYEY, PALTICK M.
REGISTRATION WIMBER: 32,411
REFERENCE/DOCKET NUMBER: PROTITELECOMMONICATION INFORMATION:
TELEPHONE: (206) 343-7074
ITELEPHONE: (206) 343-7085
INPORMATION FOR SEQ ID NO: 37: SEQUENCE CHARACTERISTICS:
LENGTH: 42 AMINO ACIDS
TYPE: AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: LINEAR
ORIGINAL SOURCE:
ORGANISM: MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 6
US-09-865-294-65
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us-09-580-018-42.rapb

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APPLICANT: Kei-Lai L. Fong C-Terminus Specific Immunosssays for TITLE OF INVENTION: N and C-Terminus Specific Immunosssays for Full Length Beta-Amyloid Peptide - Abeta(1-40), Abeta(1-39), Abeta(1-41), Abeta(1-42) and Abeta (1-43)
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100.0%; Score 217; DB 9; Length 43;
Best Local Similarity 100.4%; Pred. No. 1.66-29.

Matches 42; Conservative 0.9; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                     COUNTER READABLE FORM:

MEDIUM TYPE: 3.50 inch, 1.44MB storage
COMPUTER: IBM PC Compatibles
COMPUTER: IBM PC Compatibles
OPERATING SYSTEM: Windows
OPERATING SYSTEM: Windows
SOFTWARE: MS No. US20020182660Alepad
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/051,496
FILING DATE: 18-7a-2002
CLASSIFICATION: CURROWN-
PRICATION NUMBER: US/9/784,854A
FILING DATE: 16-Peb-2001
APPLICATION NUMBER: 60/183,407
FILING DATE: 18-Pebruary-2000
ATTORNEY/AGRAITON: AURICALIAN SAPILATION NUMBER: 60/183,407
FILING DATE: 18-Pebruary-2000
ATTORNEY/AGRAITON: AURICALIAN SAPILATION NUMBER: 60/183,407
FILING DATE: 18-Pebruary-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Koenig, C. Frederick III
REGISTRANTON UNUMER: 29,662
REFERENCE/DOCKET UNUMER: PBI-PT001.1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-6400
TELEPHONE: (215) 568-6400
                                                                                                      NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
CORRESPENCES Kei-Lai L. Fong
STREET: 1004 West 8th Avenue
CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 43 Amino Acid
TYPE: Amino Acid
STRANDEDNESS: <Unknown>
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US-10-217-459-1
; Sequence 1, Application US/10217459
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MOLECULE TYPE: Protein
HYPOTHETICAL: cUnknown>
ANTI-SENSE: cUnknown>
ORGANISM: cUnknown>
INDIVIDUAL ISOLATE: cU
CELL TYPE: cUnknown>
INDIVIDUAL ISOLATE: cU
CELL TYPE: cUnknown>
LIBRARY: cUnknown>
CLONE: cUnknown>
CLONE: cUnknown>
PUBLICATION INFORMATION:
AUTHORS:
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Patent No. US20020137112A1
GENERAL INFORMATION:
APPLICANT: Chojtker, Martina
APPLICANT: Buck, Martina
TITLE OF INVENTION: Disease
FILE REFERENCE: CHOJKIER-04302
TITLE OF INVENTION: Disease
FILE REFERENCE: CHOJKIER-04302
CURRENT APPLICATION NUMBER: US/09/731,460
CURRENT FILING DATE: 2000-12-07
NUMBER OF SEQ ID NOS: 1
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 1
LENGTH: 42
                                                         Gaps
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; Sequence 7, Application US/10076708
; Patent No. US20020164657A1
; GENERAL INFORMATION:
; APPLICANT: Sharma, Satish
; APPLICANT: Rank, Kenneth
; TITLE OF INVENTION: Assays for Accessing Aa-Tau Aggregation
; FILLE REPRENUE: 6322
; CURRENT APPLICATION NUMBER: US/10/076,708
; CURRENT FILING DATE: 2002-02-15
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 7
Query Match
100.0%; Score 217; DB 10; Length 42;
Best Local Similarity 100.0%; Pred. No. 1.5e-22;
Matches 42; Conservative 0; Mismatches 0; Indels 0
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O'THER INFORMATION: Description of Artificial Sequence:
US-09-731-460-1
                                                                                                                                          1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
                                                                                                      1 DAEFRHDSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-076-708-7
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US-09-731-460-1
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US-10-076-708-7
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Sequence 3, Application US/09808037

Sequence 3, Application US/09808037

Sequence 3. Application US/09808037

Sexent No. US20020052311A1

GENERAL INFORMATION:

METHODS METHODS AND COMPOSITIONS FOR THE TREATMENT AND/OR DIAGNOSIS OF TITLE OF INVENTION: METHODS AND DISORDERS

FILE REFRENCES SOLGMON=2D

CURRENT APPLICATION UNMERR: US/09/808,037

CURRENT FILING DATE: 2001-03-15

PRIOR FILING DATE: 2000-07-31

PRIOR FILING DATE: 1999-12-29

PRIOR FILING DATE: 1999-12-29

PRIOR FILING DATE: 1999-12-29

PRIOR FILING DATE: 1999-09-03

NUMBER OF SEQ ID NOS: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 14

US-09-304-987-1

US-09-304-987-1

Sequence 1, Application US/09904987

Patent No. US20020037908A1

GENERAL INFORMATION:

TITLE OF INVENTION: Methods and Compositions for Controlling Pathological and Prepath

TITLE OF INVENTION: Procein Assembly or Aggregation

TITLE OF INVENTION: Procein Assembly or Aggregation

FILE REFERENCE: 42108/26146

CURRENT FILING DATE: 2001-07-12

NUMBER OF SED ID NOS: 7

SOFTWARE: Patentin Version 3.0

SEQ ID NO. 1

LENGTH: 43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: homo sapiens
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: NCBI ENTREZ / QRHUA4
DATABASE ENTRY DATE: 2000-09-15
RELEVANT RESIDUES: (672)..(714)
                                                                                                                                                                                                                       MOLECULE TYPE: peptide
;
SEQUENCE DESCRIPTION: SEQ 1D NO: 1:
US-09-280-966-1
  REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-622-2300
                                                                                                INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 43 amino acids
TYPE: amino acids
TOPOLOGY: linear
                                                                             TELEFAX: 650-622-2499
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US-09-808-037-3
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                                                            APPLICANT: MUSELO, Paul
APPLICANT: MISSEN, Jeffrey
APPLICANT: MISSEN, Jeffrey
APPLICANT: TIONG, Jay
APPLICANT: TIONG, Jay
APPLICANT: TONG, JAY
APPLICANTON NUMBER: US/10/217,459
CURRENT APPLICATION NUMBER: US/10/217,459
FRIOR APPLICATION NUMBER: US/10/217,459
FRIOR PILING DATE: 1990-09-30
NUMBER OF SEQ ID NOS: 1
SOFTWARE: PATENTIN VERSION 3.0
SOFTWARE: PATENTIN VERSION 3.0
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ACID ESTERS, PHARMACEUTICAL COMPOSITIONS
COMPRISING SAME, AND METHODS FOR INHIBITING
-AMYLOID PEPTIDE RELEARE AND/OR ITS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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COMPUTER: IERS INTEST CONTROLL
CONTROLL
CONTROLL
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,966
FILING DATE: 30-Max-1999
CLASSIFICATION NUMBER: 08/976,191
FILING DATE: 21 NOV 1997
APPLICATION NUMBER: 06/077,175
FILING DATE: 22 NOV 1996
ATTORNEY/AGENT INFORMATION:
NAME: SWISS, GERALD F.
REGISTRATION NUMBER: 30,113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Burns, Doane, Swecker & Mathis, LLP
STREET: P.O. Box 1404
CITY: Alexandria
STATE: Virginia
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
; ORGANISM: beta-amyloid precursor protein
US-10-217-459-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: JAMES E. AUDIA
BEYERLY K. FOLMER
VARCHESE JOHN
JEPREY S. NISSEN
WARREN J. PORTER
EUGENE D. THORSETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/09280966
Patent No. US20010020097A1
GENERAL INFORMATION:
Publication No. US20030069445A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JING WU
                                              James
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US-09-280-966-1
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us-09-580-018-42.rapb

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0
                                                                                                                                                                          Query Match 100.0%; Score 217; DB 10; Length 43; Best Local Similarity 100.0%; Pred. No. 1.6e-22; Matches 42; Conservative 0; Mismatches 0; Indels 0; Gaps
                                                                                                                                                                                                                                                  ; SOFTWARE: Patentin version 3.0
; SEQ ID NO 3
; LENGTH: 43
; TYPE: PRT
; ORGANISM: Artificial Sequence
; PEATURE:
; PEATURE:
; OTHER INFORMATION: Synthetic peptide
US-09-808-037-3
                                                                                                                                                                                                                                                                                                                                       Search completed: April 21, 2003, 12:08:28
Job time : 15 secs
                                                                                                                                                                                                                                                      ò
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GenCore version 5.1.4 p5_4578
Copyright (c) 1993 - 2003 Compugen Ltd.
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- protein search, using sw model OM protein April 21, 2003, 12:05:20 ; Search time 18 Seconds (without alignments) 224.314 Million cell updates/sec Run on:

US-09-580-018-42

Perfect score:

1 DAEFRHDSGYEVHHOKLVFF.....DVGSNKGAIIGLMVGGVVIA 42

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

Sequence:

283224 seqs, 96134422 residues Searched:

Total number of hits gatisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

					SOMMAKIES	
Result No.	Score	* Query Match	Length	DB	ID	Description
1	217	100.0	42	10	PN0512	beta-amyloid prote
8	217	100.0	יעי	N	E60045	disea
m	217	100.0	57	~	F60045	
4	217	100.0		~	G60045	Ø
2	217	100.0	57	~	D60045	Alzheimer's diseas
9	217	100.0	ស	~	A60045	w
7	217	100.0	S	7	B60045	ò
80	217	100.0		~	PQ0438	Ø
6	217	100.0	695	7	A49795	Ø
10	217	100.0		-	QRHUA4	æ
11	198	91.2	695	7	A27485	Alzheimer's diseas
12	198	91.2		~	800550	Alzheimer's diseas
13	198	91.2		~	JH0773	Alzheimer's diseas
14	133	61.3		2	S23094	beta-amyloid prote
15	63	29.0	75	N	AI3228	tryptophan 2-monoo
16	62	28.6	75	-	QQAG4T	tryptophan 2-monoo
17	61	28.1		Н	DAAGWT	tryptophan 2-monoo
18	57	26.3	32	7	S11435	genome polyprotein
19	57	26.3	S	7	S73843	general amino acid
20	56.5	26.0	m	~	S61992	SLG1 protein - yea
21	55.5	25.6		~	G69525	formylmethanofuran
22	55.5	25.6		~	G71526	3-methyl-2-oxobuta
23	55	25.3	291	7	F95015	glycosyl transfera
24	55	25.3		7	H97888	glycosyl transfera
25	55	25.3	4	~	S27652	probable aldehyde
26	55	25.3	3063	7	JS0166	genome polyprotein
27	54.5	25.1	678	2	C81683	3-methyl-2-oxobuta
28	4.	25.1	832	~	H84848	phospholipase D [i
. 29	54	24.9	77	7	C97027	fecA-like protein,

genome polyprotein	Na+/H+ antiporter	hypothetical prote	conserved hypothet	hypothetical prote	na+/h+ antiporter	hypothetical prote	nodulin-like prote	polyprotein - pota	cation transporter	hypothetical prote	3-methyl-2-oxobuta	hypothetical prote	signal transductio	conserved hypothet	conserved hypothet
S04723	AI3443	T24727	D70128	G96774	C75103	D72302	T02581	JT0959	F86805	C86495	H72128	F70979	A81351	F70132	T18232
N	0	7	7	~	~	N	7	7	7	(7)	7	7	7	7	N
84	763	708	971	256	390	422	601	1555	314	678	678	272	339	417	527
N															
		24.7	24.7	24.4	24.4	24.4	24.4	24.4	24.2	24.2	24.2	24.0	24.0	24.0	24.0
		53.5 24.7		53 24.4			53 24.4			52.5 24.2		52 24.0	52 24.0	52 24.0	52 24.0

ALIGNMENTS

beta-amyloid protein - guinea pig (fragment) C;Species: Cavia porcellus (guinea pig) C;Date: 31-Dec-1993 #seguence_revision 31-Dec-1993 #text_change 17-Mar-1999

C) Accession: PNO512

R; Shimohigashi, Y.; Matsumoto, H.; Takano, Y.; Saito, R.; Iwata, T.; Kamiya, H.; Ohno, M. Biochem. Blophys. Res. Commun. 193, 624-630, 1993

A; Title: Receptor-mediated specific biological activity of a beta-amyloid protein fragmen A; Reference number: PNO512; MUD: 93290683; PMID: 7685598

A; Accession: PNO512

A; Accession: PNO512

A; Accession: PNO512

C; Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase ir C; Keywords: alternative splicing; amyloid

Gaps ö Length 42; Indels Query Match
Best Local Similarity 100.0%; Pred. No. 7.5e-22;
Matches 42; Conservative 0; Mismatches 0;

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RESULT 2

Alzheiner's disease amyloid beta/A4 protein precursor - sheep (fragment)
Cispecies: Ovis Sp. (sheep)
Cispecies: Ov

Length 57; 0; Indels Query Match
100.0%; Score 217; DB 2;
Best Local Similarity 100.0%; Pred. No. 1e-21;
Matches 42; Conservative 0; Mismatches 0;

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Gaps

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6 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 47 1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42 g δ

RESULT

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A,Molecule type: mRNA
A,Residues: 1-57 «JOH»
A,Cross-references: EMBL:X56125
C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase 11
C;Keywords: alternative splicing; Alzheimer's disease; amyloid; brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alzheimer's disease amyloid A4 protein precursor - rabbit (fragment)
().Species: Oryctolagus cuniculus (domestic rabbit)
().Species: Oryctolagus cuniculus (domestic rabbit)
().Species: O.Sep-1993 #sequence_revision 19-Oct-1995 #text_change 19-Oct-1995
().Accession: P00438, C60045
().Accession: P00438, C60045
().Biochem. Biophys. Res. Commun. 188, 905-911, 1992
().Aritle: Sequence and methylation in the beta/A4 region of the rabbit amyloid precursor A;Reference number: P00438 #UID:93075180; PMID:1445331
().Reference number: P00438
().Residues: 1-82 cADA>
().Residues: 1-82 cADA>
().Residues: 1-82 cADA>
().Residues: 1-82 cADA>
().Residues: No.; Changey, M.O.; Norris, P.H.; Pascual, R.; Little, S.P.
().Residues: Mol. Brain Res. 10, 299-305, 1991
().Reinerce of the Alzheimer's disease amyloid peptide in dog,
                                                                                                                                                                                                                                                                    C,Accession: A60045
R;Johnstone, E.M.; Chaney, M.O.; Norris, F.H.; Pascual, R.; Little, S.P.
Brain Res. Mol. Brain Res. 10, 299-305, 1991
A;Title: Conservation of the sequence of the Alzheimer's disease amyloid peptide in dog, A;Reference number: A60045; MUID:92017079; PMID:1656157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alzheimer's disease amyloid beta/A4 protein precursor - polar bear (fragment)
C;Species: Ursus maritimus (polar bear)
C;Species: Ursus maritimus (polar bear)
C;Species: Ursus maritimus (polar bear)
C;Species: Dec.1992 #sequence_revision 01-Dec-1992 #text_change 13-Aug-1999
C;Accession: B60045
R;Johnstone, E.M.; Chaney, M.O.; Norris, F.H.; Pascual, R.; Little, S.P.
Brain Res. Mol. Brain Res. 10, 299-305, 1991
Brain Res. Mol. Brain Res. 10, 299-305, 1991
A;Fille: Conservation of the sequence of the Alzheimer's disease amyloid peptide in of A;Reference number: A60045; MUID:92017079; PMID:1656157
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C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type ;
C;Keywords: alternative splicing; Alzheimer's disease; amyloid; brain
                                                                                                                                                                                                                        C;Species: Canis lupus familiaris (dog)
C;Date: 01-Dec-1992 #sequence_revision 01-Dec-1992 #text_change 28-Jul-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 47
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Matches 42; Conserv
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A; Residues: 1-57 <JOH>
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                                                                                       C;Date: 01-Dec-1992 #sequence_revision 01-Dec-1992 #text_change 13-Aug-1999
C;Accession: F66045
Expansion: F66045
Brain Res. Mol. Brain Res. 10, 299-305, 1991
A;Title: Conservation of the sequence of the Alzheimer's disease amyloid peptide in dog, A;Reference number: A60045; MUID:92017079; PMID:1656157
A;Mocession: F66045
A;Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Alzheimer's disease amyloid beta/A4 protein precursor - guinea pig (fragment)
C;Species: Cavia porcellus (guinea pig)
C;Species: O1-Dec-1992 #text_change 28-Jul-1995
C;Accession: G60045
R;Johnstone, E.M.; Chaney, M.O.; Norris, F.H.; Pascual, R.; Little, S.P.
Brain Res. Mol. Brain Res. 10, 299-305, 1991
A;Title: Conservation 'of the sequence of the Alzheimer's disease amyloid peptide in dog, A;Reference number:
A;Accession: G60045
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C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase
C;Keywords: alternative splicing; Alzheimer's disease; amyloid; brain
                                                                                                                                                                                                                                                                                                                                                                            A,Cross-references: EMBL:X56127; NID:g1895; PIDN:CAA39592.1; PID:g1896
C,Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase
C,Keywords: alternative splicing; Alzheimer's disease; amyloid; brain
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C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase
C;Keywords: alternative splicing; Alzheimer's disease; amyloid; brain
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C;Species: Bos primigenius taurus (cattle)
C;Date: 01-Dec-1992 #sequence_revision 01-Dec-1992 #text_change 28-Jul-1995
C;Accession: D60045
R;Johnstone, E.M.; Chaney, M.O.; Norris, F.H.; Pascual, R.; Little, S.P.
Brain Res. Mol. Brain Res. 10, 299-305, 1991
A;Title: Conservation of the sequence of the Alzheimer's disease amyloid peptide in A;Reference number: A60045; MuID:92017079; PMID:1656157
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                              .zheimer's disease amyloid beta/A4 protein precursor - pig (fragment) Species: Sus scrofa domestica (domestic pig)
Date: 01-Dec-1992 #sequence_revision 01-Dec-1992 #text_change 13-Aug-1999
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A; Residues: 1-57 <JOH>
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Residues: 1-57 <JOH>
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A; Residues: 1-57 <JOH>
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Matches 42;
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Affile: Characterization of the stend region and the first two exons of the beta-proce Affichs: Characterization of the stend region and the first two exons of the beta-proce Afficals type: DAA Accession. 12227 A3277 A3277 MUD18355979; PHID:3383813
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A;Residues: 687-622, 'G', 694-718 «XAMI»
A;Cross-references: GB:545135; NID:9557377; PIDN:AAB23645.1; PID:9257378
A;Experimental source: familial Alzheimer disease family SB
A;Notes: sequence extracted from NCBI backbone (NCBIP:115374)
A;Accession: B44017
A;Molecule type: DNA
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Ni Container, s disease amyloid beta protein precursor [validated] - human

Ni Alternate names. Alzienimer's disease amyloid Af protein; coagulation factor XIa inhibi

Ni Contains amyloid beta protein long, plaque form; amyloid beta protein short, vascular

protein precursor splice form APP(770)

C; Specials: Hono saples (man)

C; Date: 30-Jun-1887 Hesquence revision 28-Jul-1895 #text change 15-Sep-2000

C; Date: 30-Jun-1887 Hesquence revision 28-Jul-1895 #text change 15-Sep-2000

C; Date: 30-Jun-1887 Hesquence revision 28-Jul-1895 #text change 15-Sep-2000

C; Date: 30-Jun-1887 Hesquence revision 28-Jul-1895 #text change 15-Sep-2000

C; Date: 30-Jun-1887 Hesquence revision 28-Jul-1895 #text change 15-Sep-2000

C; Date: 30-Jun-1887 Hesquence revision 28-Jul-1895 #text change 15-Sep-2000

A; Molecule A, 38-Sep 31, A29302; A60895; Jul-1895 A, 37-Sep 31, A38-384; Jul-1895 A, 38-Sep 31, 38-Sep 31, A38-384; Jul-1895 A, 38-Sep 31, A38-384; Jul-1895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Reference number: A60045; MUID:92017079; PMID:1656157
A;Aoceaule type: mRNA
A;Molecule type: mRNA
A;Residues: 12-68 < UOH>
A;Crose-references: MBNB:X56129
C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase C;Keywords: alternative splicing; Alzheimer's disease; amyloid; Down's syndrome
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es 42; Conservative
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Cross-references: GB:X06981; NID:928816; PIDN:CAA30041.1; PID:9929611
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A;Molecule type: DNA
A;Residues: 1-19 <RES>
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A; Recessions: 254-288, VV, 365-646, E', 648-770 < ROB>
A; Molecule type: mRNA
A; Residues: 254-288, VV, 365-646, E', 648-770 < ROB>
A; A; Cross-references: GB:MiG-55; NID:9178539; PIDN:AAA51722.1; PID:9178540
A; A; Cross-references: GB:MiG-56; NID:9178539; PIDN:AAA51722.1; PID:9178540
B; Goldeaber; D: Lesman, M.1:) McBride, O.W.; Saffiotti, U.; Gajdusek, D.C.
A; Title: Chisacterization and chromosomal localization of a cDNA encoding brain amyloid
A; Reference number: A47584; MUD:81720328; PMID:3810169
A; Accession: A47584; MUD:81720328; PMID:3810169
A; Accession: A47584; MUD:81720328; PMID:3810169
A; Reference number: A47584; MUD:8178706; PIDN:AAA35540.1; PID:9178707
A; Reference S; GB:Mi5533; NID:9178706; PIDN:AAA35540.1; PID:917870
A; Reference S; GB:Mi5533; NID:9178706; PIDN:AAA35540.1; PID:9177958
A; Tatle: A47584; MUD:81720329; PMID:2949367
A; Title: A47585; MUD:87120329; PMID:2949367
A; Reference number: A47585
A; Molecule type: mRNA
A; References: GB:Mi5532; NID:9177957; PIDN:AAA51564.1; PID:9177958
A; Molecule type: mRNA
A; Reference number: S02638; MUD:89249367
A; Miccession: S02638; MUD:892949367
A; Molecule type: mRNA
A; Reference number: S02638; MUD:89294937; PMID:2949367
A; Miccession: S02638; MUD:89296437; PMID:2949367
A; Molecule type: mRNA
A; Reference number: S02638; MUD:89296437; PMID:290137
A; Reference number: S02638; MUD:89296437; PMID:290137
A; Reference number: S02638; MUD:89296437; PMID:290137
A; Reference number: S02638; MUD:8928437; PMID:29093290
A; Reference number: S02638; MUD:8928437; PMID:29093290
A; Reference number: S02638; MUD:89282640; PMID:2893290
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Residues: 286-344'.1',365-366 <TAN2>
Residues: 286-344'.1',365-366 <TAN2>
Fresidues: 286-344'.1',365-366 <TAN2>
Fresidues: 286-344'.1',365-366 <TAN2>
Fresidues: 286-344'.1',365-366 <TAN2>
Fresidues: 286-34'.1',365-36'.1'
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Fresidues: 286-36'.1'
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FResidues 1.34,71,365-770 <PO2>
FResidues 1.34,71,365-770 <PO2>
Cross-references: GB:X06989; EMBL:Y00397; NID:928720; PIDN:CAA30050.1; PID:928721
Note: alternative splice form APP(751)
Kitaganchi, N. 7 Takahashi, Y.; Tokushima, Y.; Shiojiri, S.; Ito, H.
Ature 331, 530-532, 1988
Tifile: Novel precursor of Alzheimer's disease amyloid protein shows protease inhibitor
AREference number: A38949; MUID:88122641; PMID:2893291
Regidues: 687-718 «KAM2»

(Zroser-references GB: 5843.36; NID:9257379; PIDN:AAB23646.1; PID:9257380

(Zroser-references GB: 5843.36; NID:9257379; PIDN:AAB23646.1; PID:9257380

(Skaperimental source: familial Alzheimer disease family LIT

(Note: sequence extracted from NCBI backbone (NCBIP:115376)

(Note: this sequence has a silent muration

(Kang, J.; Lemaire, H.G.; Unterbeck, A.; Salbaum, J.M.; Masters, C.L.; Grzeschik, K.H.;

(Skap, J.; Lemaire, H.G.; Unterbeck, A.; Salbaum, J.M.; Masters, C.L.; Grzeschik, K.H.;

(Title: The precursor of Alzheimer's Glassase amyloid A4 protein resembles a cell-surfac
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Residues 1-288, V. 365-770 «KAN»

(Tross-references: GB:Y00264; NID:g28525; PIDN:CAA68374.1; PID:g28526

(Note: alternative splice form APP(655)

(Robakia, N.K.; Ramakrishna, N.; Wolfe, G.; Wisniewski, H.M.

FRObakia, N.K.; Ramakrishna, N.; Wolfe, G.; Wisniewski, H.M.

Froc. Natl. Acad. Sci. U.S.A. 84, 4190-4194, 1987

Title: Wolecular cloning and characterization of a cDNA encoding the cerebrovascular increasion: A29030, MUID:87231971; PMID:303574
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Residues: 287-367 <KIT>
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A, Experimental source: glioblastoma cell line
A, Experimental source: glioblastoma cell line
A, Experimental source: glioblastoma cell line
B, Wittek, M.P.; Rasool, C.G.; de Sauvage, F.; Vitek, S.M.; Bartus, R.T.; Beer, B.; Ashton
B, Vitek, M.P.; Rasool, C.G.; de Sauvage, F.; Vitek, S.M.; Bartus, R.T.; Beer, B.; Ashton
B, Vitek, M.P.; Rasool, C.G.; de Sauvage, F.; Vitek, S.M.; Bartus, R.T.; Beer, B.; Ashton
A, Reference number: A30320
A, Reference number: A30320
A, Status: not compared with conceptual translation
A, Molecule type: mRNA
A, Residues: 284-288, V., 365-770 «VIT1>
A, Accession: B30320
A, Status: not compared with conceptual translation
A, Molecule type: mRNA
A, Residues: c30320
A, Status: not compared with conceptual translation
A, Molecule type: mRNA
A, Residues: not compared with conceptual translation
A, Molecule type: mRNA
A, Residues: c30320
A, Status: not compared with conceptual translation
A, Molecule type: mRNA
A, Residues: c30320
A, Status: not compared with conceptual translation
A, Molecule type: mRNA
A, Residues: c30320
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A, Molecule type: mRNA
A, Residues: c30320
A, Status: not compared with conceptual translation
A, Molecule type: mRNA
A, Residues: c30320
A, Status: not compared with conceptual translation
A, Molecule type: mRNA
A, Residues: c60-770 «VIT3>
A, Residues: 607-770 «AIT3>
A, Residues: 507-770 «AIT3>
A, Residues: 507-770 «AIN3
A, Molecule type: mRNA
A, Residues: 607-770 «AIN3
A, Molecule type: mRNA
A, G, G, G, C, Residue 609 as Asn, 
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Alzheimer's disease amyloid beta/A4 protein homolog precursor - mouse
NiAlternate names: proteinase nexin II
CiSpecias: Mus musculus (house mouse)
CiDate: 31-Mar-1989 #sequence revision 31-Mar-1989 #text_change 13-Aug-1999
CiAccession: A27485, S19727; I49485
R;Yamada T.; Saakii, H; Puruya, H; Miyata, T.; Goto, I.; Sakaki, Y.
Biochem Biophys. Res. Commun. 149, 665-671, 1987
A;Title: Complementary DNA for the mouse homolog of the human amyloid beta protein precur A;Meference number: A27485; MUD:88106489; PMID:3322280
A;Mocession: A27485
A;Accession: A27485
A;Accession: A27485
A;Accession: D-695 c/AMA
A;Reference number: A27485; MUD:9191568; PIDN:AA37139.1; PID:9309085
A;Accession: Bource: brain
B;Accession: 
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Arryckopan 2-monooxygenase tms1 [imported] - Agrobacterium tumefaciens (strain C59, Dupor C5, Decides Agrobacterium tumefaciens
C;Decides Agrobacterium tumefaciens
C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 01-Feb-2002
C;Accession: A13228
A;Mood, Duy, Setublal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Mood, G.E.; Chen, X.; Moo, L. R;Mood, O.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClells
A;Arzp, P.; Romero, P.; Zhang, S.
Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm, F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 14

52,3094

beta-amyloid protein precursor - rat

C;Species: Rattus norregicus (Norway rat)

C;Species: Rattus norregicus (Norway rat)

C;Accession: 823094

C;Accession: 823094

A;Title: Two-way cleavage of beta-amyloid protein precursor by multicatalytic proteinase

A;Title: Two-way cleavage of beta-amyloid protein precursor by continated by A;Accession: 823094

A;Molecule type: protein

A;Molecule type: protein

A;Residues: 1-33 - KOJ>

C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase 11
R;Okado, H.; Okamoto, H. Bes. Commun. 189, 1561-1569, 1992
blochem. Balphys. Res. Commun. 189, 1561-1569, 1992
A;Title: A Xenopus homologue of the human beta-amyloid precursor protein: developmental a A;Reference number. JH0773; MUID:9312927; PMID:1282805
A;Accession. JH0773
A;Accession. JH0773
A;Residues: 1.747 < OKA>
A;Residues: 1.747 < OKA>
A;Coss_references GB:552417; NID:9263150; PIDN:AAB24853.1; PID:9263151
A;Experimental source: larva disease amyloid beta protein; animal Kunitz-type proteinase in C;Keywords: alternative splicing; amyloid
F;287-337/bomain: animal Kunitz-type proteinase inhibitor homology <BPI>
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A;Accession: A13228
A;Accession: A13228
A;Accession: A13228
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-755 KUR>
A;Ceces-references: GB:AB008690; PIDN:AAL46247.1; PID:g17744025; GSPDB:GN00189
A;Crose-references: GB:AB008690; BDN:AAL46247.1; PID:g17744025; GSPDB:GN00189
A;Experimental source: strain C58 (Dupont)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ő
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Best Local Similarity 89.3%; Pred. No. 6.6e-11;
Matches 25; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 91.2%; Score 198; DB 2; Length 74°
Best Local Similarity 88.1%; Pred. No. 5.5e-18;
Matches 37; Conservative 4; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      649 DSEYRHDTAYEVHHQKLVFFAEEVGSNKGAIIGLMVGGVVIA 690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 DAEFRHDSGYEVHHOKLVFFAEDVGSNKGALIGLMVGGVVIA 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 DAEFGHDSGFEVRHQKLVFFAEDVGSNK 33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alzheimer's disease amyloid beta protein precursor - rat
Nylhernate names: beta-A4 amyloid protein
Nylhernate names: beta-A4 amyloid protein
C;Species; Rattus norvegicus (Norway rat)
C;Species; Rattus norvegicus (Norway rat)
C;Species: Taliniu-1989 #sequence revision 30-Jun-1989 #text_change 13-Aug-1999
C;Accession: S00550; A41245; Ā39920; S46251
R;Shivers, B.D.; Hillich, C.; Multhaup, G.; Salbaum, M.; Beyreuther, K.; Seeburg, P.H.
R;Shivers, B.D.; Hillich, C.; Multhaup, G.; Salbaum, M.; Beyreuther, K.; Seeburg, P.H.
R;ATitle: Alzheimer's disease amyloidogenic glycoprotein: expression pattern in rat brain
A;Reference number: S00550; MUD:88312583; PMID:2900758
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Molecule type: BMBL:X07648; NID:955616; PIDN:CAa30488.1; PID:955617
R;Schubert, D.; Schroeder, R.; LaCorblere, M.; Saltoh, T.; Cole, G.
S;Gence 241, 223-226, 1988
A;Reference number: BMLD:355 MUD:88264430; PMID:2968652
A;Accession: A41245; MUD:88264430; PMID:2968652
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Experimental source: brain content as neurofibrillary tangles and/or plaques is content. Deposition of amyloid protein as neurofibrillary tangles and/or plaques is comperfamily. Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase feepwords: alternative splicing, amyloid, glycoprotein; transmembrane protein 625-648/Domain: transmembrane #status predicted <TWM>
    A,Cross-references: GB:D10603; NID:g220328; PIDN:BAA01456.1; PID:g220329
C,Genetics:
A,Map position: 16C3
C;Superfamily: Alzheimer's disease amyloid beta protein; animal Kunitz-type proteinase C;Keywords: alternative splicing; amyloid; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                           Length 695;
                                                                                                                                                                                                       Query Match 91.2%; Score 188; DB 2; Length 69
Best Local Similarity 92.9%; Pred. No. 5.1e-18;
Matches 39; Conservative 1; Mismatches 2; Indels
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A, Molecule type: protein
A, Molecule type: protein
A, Readidues: 18-32 < POT.
A, Experimental source: brai
C, Comment: Deposition of am
C, Superfamly: Alzheimer's
C, Keywords: alternative spl
F, 625-648/Domain: transmemb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
es 39; Conserv
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Matches
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A;Genome: plasmid C;Superfamily: Agrobacterium plasmid tryptophan 2-monooxygenase

Score 63; DB 2; Pred. No. 3.3;

29.08;

Query Match Best Local Similarity

RESULT 13
JHO771

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6; Gaps 9; Indels Matches 16; Conservative 5; Mismatches

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Search completed: April 21, 2003, 12:07:46 Job time : 20 secs

us-09-580-018-42.rsp

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April 21, 2003, 12:01:04 ; Search time 11 Seconds (without alignments) 158.364 Million cell updates/sec
                                                                                                                                                                        US-09-580-018-42
217
1 DAEFRHDSGYEVHHQKLVFF.....DVGSNKGALIGLMVGGVVIA 42
                                                                                                                                                                                                                                                                                                                                      112892
GenCore version 5.1.4 p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                112892 seqs, 41476328 residues
                                                                                                                                                                                                                                                                                                                                                                                                                          Post-processing: Minifum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                  OM protein - protein search, using sw model
                                                                                                                                                                                                                                              BLOSUM62
Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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                                                                                                                                                                          Scoring table:
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                                                                                                          Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No. 8	Score	% Query Match	Length	DB	SUMMAKIES	Description	
	217	100.0	57	-	A4 PIG	Q29023 sus scrofa	,
N	217	100.0		٦		ursus	ц
m	217	100.0	58	٦	A4 CANFA		_
4	217	100.0		٦	A4_RABIT	-	œ
z,	217	100.0	Ŋ	-	A4_SHEEP	Q28757 ovis aries	
9	217	100.0		Н	A4_BOVIN		
7	217	100.0	7	Н	A4 SAISC		
æ	217	100.0		Н	A4_HUMAN	P05067 homo sapien	ď
6	198	91.2		Н	A4_MOUSE		3
0	198	91.2	770	Н	A4_RAT		>
_	62	28.6	755	~	TRZM AGRT4	P04029 agrobacteri	·~
7	61	28.1		Н	TR2N_AGRVI		٠,۲
٣	57	26.3	327	Н	POLG_PVYCH		2
4	57	26.3		Н	Y226_MYCPN	P75462 mycoplasma	
S	56.5	26.0		Н	SLG1_YEAST		υ
9		25.6	297	н	FTR ARCFU		Q
7	55	25.3	488	Н	DHAL PSESP		ĽΩ
80	55	25.3	٣	~	POLG_PVYN		0
6	54.5	25.1	196	~	PDB1_ARATH		ω
20	54	24.9	284	Н	POLG_PVYYO	P11897 potato viru	n
4	53.5	24.7	708	Н	YNZB_CAEEL	-	٠,
22	53.5	24.7	971	7	Y228_BORBU	_	n
6	25	24.0	611	~	YCR3_YEAST	P25351 saccharomyc	ט
4	51	23.5		~	COBQ_MYCTU	_	٠,-
'n	21	23.		٦	ITB2 BOVIN		
26	50.5	23.		7	PDB2_ARATH		ĽΩ
27	20	23.	285	-	MENB HAEIN	P44960 haemophilus	D
28	50			~	MRP5 HUMAN		ㅁ
29	20			7	STAN DROME	_	
0	49.5	22.8	٦		VGL2_IBVM	_	υ
7	49	22.6		-	TPIS_CULTA	_	ಹ
7	49	22.6			Y853 RICPR	Q92ca7 rickettsia	
۳	4	22.6		Н	PGK STRCO	Q9z519 streptomyce	ø

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01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
01-NOV-1997 (Rel. 40, Last annotation update)
Alzheimer's disease amyloid A4 protein homolog [Contains: Beta-amyloid protein (Beta-APP) (A-beta)] (Fragment)
Alzheimer's disease amyloid A4 protein homolog (Contains: Beta-amyloid protein (Beta-APP) (A-beta)] (Fragment).
                                                                                                                                                   TISSUE=Kidney,
MEDLINE=2017079; PubMed=1656157;
MEDLINE=20210709; PubMed=1656157;
MEDLINE=20210709; PubMed=1656157;
MEDLINE=20210709; PubMed=1656157;
MCOnservation of the sequence of the Alzheimer's disease amyloid peptide in dog, polar bear and five other mammals by cross-species polymerase chain reaction analysis.";
Brain Res. Mol. Brain Res. 10:299-30511991.
INTRACELLULAR SIGNALING PATHMAY THROUGH THE GTP-BINDING PROTEIN GIO) INTRACELLULAR SIGNALING PATHMAY THROUGH THE GTP-BINDING PROTEIN -:- SUMCEBLULAR LOCATION: TYPE I membrane protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Brain;
MEDILNE-92017079; PubMed=1656157;
Johnstone E.M., Chaney M.O., Norris F.H., Pascual R., Little S.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus
                                              caris familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BETA-AMYLOID PROTEIN (POTENTIAL).
EXTRACELLULAR (POTENTIAL).
POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 217; DB 1; Length 58; Similarity 100.0%; Pred. No. 8.1e-22; Conservative 0; Mismatches 0; Indels 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    49 BETA-AMYLOID PROTEIN (PC
34 EXTRACELLULAR (POTENTIAL
58 POTENTIAL.
58
6285 MW; 8469D488A2E12DFA CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR001868; A4_APP.
InterPro; IPR001255; Beta-APP.
PR031494; Beta-APP; 1.
PR05ITE; PS00319; A4_EXTRA; PARTIAL.
PR0SITE; PS00310; A4_INTRA; PARTIAL.
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                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        58 AA;
                                                                                                   NCBL TaxID=9615;
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Local Simi
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Q28748;
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SEQUENCE
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                                                                                                                A4_URSMA STANDARD; PRT; 57 AA.

Q29149;

Q1-NOV-1997 (Rel. 35, Created)

10-NOV-1997 (Rel. 35, Last sequence update)

30-MAY-2000 (Rel. 39, Last annotation update)

Alzhaimer's disease amvioid A4 protein homolog [Contains: Beta-amyloid protein (Beta-APP) (A-beta)] (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                          Ursus maritimus (Polar bear) (Thalarctos maritimus).
Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Fissipedia; Ursidae; Ursus.
NCBI_TAXID=29073;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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EXTRACELLULAR (POTENTIAL).
POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84209D88EBA82DFA CRC64;
6 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -1- SUBCELLULAR LOCATION: Type I membrane protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 217; DB 1;
Pred. No. 7.9e-22;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HSSP; P05067; 1BA4.
INCEPTO; IPR01868; A4_APP.
INCEPTO; IPR001255; Beta-APP.
Fam; PF03494; Beta-APP; PROSITE; PS00319; A4_EXTRA; PARTIAL.
PROSITE; PS00319; A4_EXTRA; PARTIAL.
PROSITE; PS00320; A4_INFRA; PARTIAL.
PNON TER
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(Rel. 35, Last sequence update)
(Rel. 39, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      58 AA.
                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=92017079; PubMed=1656157;
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ilarity 100.0%;
Conservative 0;
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SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            57 AA;
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Gaps

01-NOV-1997 (01-NOV-1997 (30-MAY-2000 (

A4 CANFA Q28280;

RESULT 3
A4 CANPA
ID A4 C
AC
AC 0282:
DT 01-N
DT 30-M

42;

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DOMAIN TRANSMEM NON TER

CHAIN

SEQUENCE Query Match Local Best Loca Matches

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01-NOV-1997 (Rel. 35, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
41rhaimer's disease amyloid A4 protein homolog [Contains: Beta-amyloid protein (Beta-APP) (A-beta)] (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Brain;
MEDLINE=2010709; PubMed=1656157;
MEDLINE=2010709; PubMed=1656157;
MEDLINE=2010709; PubMed=1656157;
MEDLINE=2010709; PubMed=1656157;
MEDLINE=2010709; PubMed=1656157;
MEDLINE=2010709; PubMed=1656159;
MOLES MOLE
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BETA-AMYLOID PROTEIN (POTENTIAL). EXTRACELLULAR (POTENTIAL).
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CYTOPLASMIC (POTENTIAL).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INCEPTCO 18001868; A4_APP.
INCEPTCO 18001825; Beta-APP.
Fram; PF03494; Beta-APP; 1.
PROSITE; PS00319; A4_EXTRA; PARTIAL.
PROSITE; PS00320; A4_INTRA; PARTIAL.
GLYCOPTCCEIN; AM, 101 di Neurone; Transmembrane.
NON TER
CHAIN
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100.0%; Pred. No. 8.1e-22;
iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; X56130; CAA39595.1; -. HSSP; P05067; 1BA4.
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Q28053;
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NON TER
SEQUENCE
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Matches
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A4_BOVIN
                       DAY WAREN TO COLOR COLOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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01-NOV-1997 (Rel. 35, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Alzheimer's disease amyloid A4 protein homolog [Contains: Beta-amyloid protein (Beta-APP) (A-beta)] (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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WEDLINES-2017079; PubMed=1656157;

WE
     "Conservation of the sequence of the Alzheimer's disease amyloid peptide in dog, polar bear and five other mammals by cross-species polymerase chain reaction analysis.";

Brain Res. Mol. Brain Res. 10:299-305(1991).

--- FUNCTION: FUNCTIONAL NEURONAL RECEPTOR WHICH COUPLES TO INTRACELLULAR SIGNALING PATHWAY THROUGH THE GTP-BINDING PROTEIN G(O) [BY SIMILARITY).

--- SUBCELLULAR LOCATION: Type I membrane protein.
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Bukaryota, Metazoa; Chordata, Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria, Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Caprinae; Ovis.
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; Pred. No. 8.1e-22;
0; Mismatches 0; Indels
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NON TER 1 1
CHAIN 6 48 BETA-AMYLOID PROT
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Interpro, IRR001255; Beta-APP.
Edm, PF03494; Beta-APP; 1.
PROSITE; PS00319; A4_EXTRA, PARTIAL.
PROSITE; PS00320; A4_INTRA; PARTIAL.
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6300 MW;
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nes 42; Conservative
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Q28757;
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A4 SHEEP

10 A4 SHEEP

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SEQUENCE FROM N.A.
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-!- DOMAIN: THE CLATHRIN-BINDING SITE IS ESSENTIAL FOR ITS ASSOCIATION
WITH XII ALPHA, -BETA, AND -CAMAA THE SEQUENCE SPECIFIC
RECOGNITION EXTENDS TO PEPTIDE RESIDUES THAT ARE C-TERMINAL TO THE
NPXY MOTIF. THIS INTERACTION APPEARS TO BE INDEPENDENT OF
PHOSPHORYLATION (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE APP FAMILY.
-!- SIMILARITY: CONTAINS 1 BPII/KUNITZ INHIBITOR DOMAIN.
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Eukaryota, Metazoa, Chordata, Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini, Cebidae; Cebinae; Saimiri.
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15-DBC-1998 (Rel. 37, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)

Alzheimer's disease amyloid A4 protein precursor [Contains: Beta-amyloid protein (Beta-APP) (A-beta)].
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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HSSP, P05067; 1BA4.

InterPro; IPR001255; Beta-APP.
InterPro; IPR001255; Beta-APP.
Pfam; PP03494; Beta-APP; 1.
PROSITE; PS00319; A4 EXTRA; PARTIAL.
PROSITE; PS00320; A4 INTRA; PARTIAL.
GlyCoprotein; Amyloid; Neurone; Transmembrane.
NON TER 7 49 BETA-AMYLOID PROTEIN
DOWAIN <1 34 EXTRACELLULAR (POTEN
TRANSMEM 35 58 POTENTIAL.
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Pred. No. 8.2e-22;
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llarity 100.0%; Pred. No. 8.2
Conservative 0; Mismatches
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InterPro; IPR001255; Beta-APP.
InterPro; IPR00223; Kunitz BPTI.
Pfam; PF00117; A4_EXTRA; 1.
                                                                                                                                                                                                                                                                                                                                                                               EMBL; S81024; AAD14347.1; -. HSSP; P05067; 1AAP.
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Local Sim-
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Q95241;
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SEQUENCE
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Matches
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A4-SAISC

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P05067; P09000; Q16011;
13.40G-1997 (Rel. 05, Created)
01-NOV-1991 (Rel. 20, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Alzheimer's disease amyloid A4 protein precursor (Protease nexin-II)
(PN-II) (Contains: Beta-amyloid protein (Beta-APP) (A-beta)].
APP OR A4 OR CVAP OR AD1.
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kang J., Lemaire H.-G., Unterbeck A., Salbaum J.M., Masters C.L., Grzeschik K.-H., Multhaup G., Beyreuther K., Mueller-Hill B.; "The precursor of Alzheimer's disease amyloid A4 protein resembles cell-surface receptor.";
Nature 325:733-736(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryoča; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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MEDLINE-89128427; PubMed-2783775;
Lemaire H.G., Salbaum J.M., Multhaup G., Kang J., Bayney R.M.,
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BEDILNES-BRIZZ639; PubMed=2893289;

PORTE P., GORZA-BR-DeWhitt P., Schilling J., Miller J., HSu D.

Greenberg B., Davis K., Wallace W., Lieberburg I., Fuller F.,
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MEDLINE=87144572; PubMed=2881207;
PRINTS; PR00203; AMYLOIDA4.
PRINTS; PR00759; BASICPTASE.
ProDom; PD000222; Kunitz_BP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     proteinase inhibitors.";
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and partial sequence of a 4,200-dalton peptide isolated from cortical Unterbeck A., Beyreuther K., Mueller-Hill B.; "The PreA4(655) precursor protein of Alzheimer's disease A4 amyloid is encoded by 16 exons."; "is encoded by 17.517-522(1989). for SEQUENCE OF 286-345 AND 365-366 FROM N.A.
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"Purification of protease nexin II from human fibroblasts.";
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COMPLEX WITH G(O).

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Murayama Y., Ogata E.;

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IDENTITY OF APP WITH NEXIN-II.
MEDLINE-89384866; PubMed=2506449;
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Johnson-Wood K.L., Beattie E.C., Ward P.J., Blacher R.W., Dovey H.F.,
Sinha S.;
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"Three-dimensional structures of the amyloid beta peptide (25-35) in
membrane-mimicking environment.";
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Kamarck M.E., Ramabhadran T.V., Dreyer R.N., Davis G., Unterbeck A.,
Tamburini P.P.,
"Sequential P.P.," Sequence assignment and structure determination of
the Kunitz-type inhibitor domain of the Alzheimer's beta-amyloid
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                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=90211252; PubMed=1969731; Kido H., Fukutomi A., Schilling J., Wang Y., Cordell B., Katunuma "Protease-specificity of Kunitz inhibitor domain of Alzheimer's
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Galatis D., Hesse L., Multhaup G., Beyreuther K., Masters C.L.,
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domain of
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STRUCTURE BY NMR OF 672-699.
MEDLINE=20400066; PubMed=10940222;
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Biochemistry 35:16094-16104(1996)
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                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Brain;
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Yamada T., Sasaki H., Furuya H., Miyata T., Goto I., Sakaki Y.;
"Complementary DNA for the mouse homolog of the human amyloid beta
                                                                                                                                                                [24]
SIGNAL SEQUENCE CLEAVAGE SITE, AND TOPOLOGY.
SIGNAL SEQUENCE CLEAVAGE SITE, AND TOPOLOGY.
BYEDLINE=88296437; PubMed=2900137;
Dyrks T., Waldeman A., Multhaup G., Salbaum J.M., Lemaire H.-G.,
Kang J., Mueller-Hill B., Masters C.L., Beyreuther K.;
"Identification, transmembrane orientation and biogenesis of the amyloid A4 precursor of Alzheimer's disease.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI TaxID=10090;
            "Solution structures in aqueous SDS micelles of two amyloid beta peptides of Abeta(1-28) mutated at the alpha-secretase cleavage
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                     01-OCT-1989 (Rel. 12, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
16-OCT-2001 (Rel. 40, Last annocation update)
Alzhaimer's disease amyloid A4 protein homolog precursor (Amyloidogenic glycoprotein) (AG).
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Submitted (MAR-1988) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                      672 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 713
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Biochem. Biophys. Res. Commun. 149:665-671(1987)
                                                                                                                                                                                                                                                                                                                                                                                                              770 AA.
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Poulsen S.-A., Watson A.A., Craik D.J.
                                                  Struct. Biol. 130:142-152(2000).
                                                                                                                                                      Struct. Biol. 130:130-141(2000).
                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
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Exercise 2019955; publications of the Sackit H., Hattori M., Randa T., Yoshikai S.I., Sasaki H., Hattori M., Sakii Y.; Yamada T., Yoshikai S.I., Sasaki H., Hattori M., AND ALTERNATIVE SPLICING.

R. Althemer's disease amyloid precursor-encoding gene in mouse.'; density 12:199-195(192).

R. Althemer's disease amyloid precursor-encoding gene in mouse.'; density 13:199-195(192).

R. Althemer's disease amyloid precursor-encoding gene in mouse.'; density 13:199-195(192).

R. Yamada T., Sasaki H., Dohura K., AND ALTERNATIVE SPLICING.

R. TISSUB-SERIES BRIDGE OF Althemyl. AND ALTERNATIVE SPLICING.

R. TISSUB-SERIES BRECHFORNERS BR. AND ALTERNATIVE SPLICING.

R. ALTERNATIVE RENOWERS Res. Commun. 16:196-191(196).

R. ALTERNATIVE RENOWERS Res. Commun. 16:196-191(196).

R. ALTERNATIVE RENOWERS Res. AND CAMPAR APP (195), APP (
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SEQUENCE FROM N.A.
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TR2M AGRT4
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SIMILARITY: BELONGS TO THE APP FAMILY.
SIMILARITY: CONTAINS I BPTI/KUNITZ INHIBITOR DOMAIN.
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Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus
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"The sequence of the two extra exons in rat preA4.";
Nucleic Acids Res. 17:2130-2130(1989).
-!- SUBCELLULAR LOCATION: Type I membrane protein.
-!- ALTERNATIVE PRODUCTS: 5 ISOFORMS; APP(395), APP(563), APP(751) AND APP(770) (SHOWN HERE); ARE PRODUCED BY ALTERNATIVE SPLICING.
                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Brain;
MIDLINE=88312583; PubMed=2900758;
Shivers B.D., Hilbich C., Multhaup G., Salbaum J.M., Beyreuther K., Seeburg P.H.; Harshaimer's disease amyloidogenic glycoprotein: expression pattern in rat brain suggests a role in cell contact.";
EMBO J. 7:1365-1370(1988).
BPTI/KUNITZ INHIBITOR.
CLATHRIN-BINDING (BY SIMILARITY).
BY SIMILARITY.
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BY SIMILARITY.
N-LINKED (GLCNAC. . ) (POTENTIAL).
N-LINKED (GLCNAC. . ) (POTENTIAL).
E -> V (IN ISOFORM APP(655)).
MISSING (IN ISOFORM APP(655)).
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MISSING (IN ISOFORM APP(551)).
W; 26CSODE0890CAF7A CRC64;
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Pred. No. 3.4e-18;
1; Mismatches 2; Indels
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01-DEC-1992 (Rel. 24, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Alzheimer's disease amyloid A4 protein homolog precursor (Amyloidogenic glycoprotein) (AG),
                                                                                                                                                                                                      672 DAEPGHDSGFEVRHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 713
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MEDLINE-89183625; Pubmed=2648331;
                                                                                                                     86752 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE OF 289-364 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; X07648; CAA30488.1; -. EMBL; X14066; CAA32229.1; -.
                                                                                                                                           91.2%;
                                                                                                                                                                   Matches 39; Conservative
                                                                                                                                                                                                                                                                          STANDARD;
                                                                                                                                                                                                                                                                                                                                                                      (Rat)
                                                                                                                     770 AA;
                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                       Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10116;
  287
759
300
316
542
571
289
346
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VARSPLIC
VARSPLIC
SEQUENCE
               SITE
DISULFID
DISULFID
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                                                                                                                                             Query Match
                                                           CARBOHYD
                                                                                                                                                                                                                                                                         A4 RAT
P08592;
  DOMAIN
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SEQUENCE FROM N.A.
STRAIN-Ach5; PLASMID=PTIAch5;
MEDLINE=84207942; PubMed=6327292;
MEDLINE=84207942; PubMed=6327292;
Gielen J., de Beuckeleer M., Seurinck J., Deboeck F., de Greve H.,
Lemmers M., van Montagu M., Schell J.;
"The complete nucleotide sequence of the TL-DNA of the Agrobacterium
tunefaciens plasmid PTIAch5.";
EMBO J. 3:835-846(1984)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ALZHEIMER'S DISEASE AMYLOID A4 PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CYTOPLASMIC (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
EQUIVALENT OF BETA-AMYLOID PROTEIN.
BPTI/KUNITZ INHIBITOR.
CLATHRIA-BINDING (BY SIMILARITY).
BY SIMILARITY.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Agrobacterium tumefaciens.
Plasmid priA6NC.
Bacteria, Proteobacteria; alpha subdivision; Rhizobiaceae group;
Rhizobiaceae; Rhizobium.
NCBI_TaxID=176298, 358;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-LINKED (GLCNAC. . .) (POTENTIAL)
N-LINKED (GLCNAC. . .) (POTENTIAL)
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E -> V (IN ISOFORM APP(695)).
MISSING (IN ISOFORM APP(695)).
C26C9DGBEDD29A7 (CKC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 770;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         91.2%; Score 198; DB 1; Length 77 92.9%; Pred. No. 3.4e-18; ive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                  PROSITE; PS00319; A4_EXTRA; 1.
PROSITE; PS00320; A4_INTRA; 1.
PROSITE; PS00280; BPTI KUNITZ 1; 1.
PROSITE; PS50279; BPTI_KUNITZ 2; 1.
Glycoprotein; Amyloid; Neurone; Transmembrane; Signal; Allernative splicing; Serine procease inhibitor.
SIGNAL 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HOMOLOG.
EXTRACELLULAR (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       713
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    672 DAEFGHDSGFEVRHOKLVFFAEDVGSNKGAIIGLMVGGVVIA
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23-0CT-1986 (Rel. 02, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Iryptophan 2-monooxygenase (EC 1.13.12.3).
86704 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 92.9 Matches 39; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
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770 AA;
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RESULT 13
POLG PVYCH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ij
                                      Fuller S., Montoya A., Horodoyski P., Lichtenstein C., Garfinkel D., Fuller S., Flores C., Peschon J., Nester E., Gordon M.;
Fuller S., Flores C., Peschon J., Nester E., Gordon M.;
"Nuclectide sequence of the tama genes of the Pilakoc octopine Ti
plasmid: two gene products involved in plant tumorigenesis.";
Proc. Natl. Acad. Sci. U.S.A. 81:1728-1732(1984).
-!- CAPALYITC ACTIVITY: L-tryptophan + O(2) = indole-3-acetamide +
CO(2) + H(2)O.
-!- PATHWAY: FIRST STEP IN THE BIOSYNTHESIS OF AUXINS FROM TRYPTOPHAN.
-!- MISCELLANBOUS: THE SEQUENCE SHOWN IS TEAT OF ACHS.
-!- SIMILARITY: SIGNIFICANT HOMOLOGY TO THE ADENINE BINDING REGION OF
P-HYDROXYBENZOATE HYDROXYLABE FROM P. FLUORESCENS. IT SEEMS THAT
THIS PROTEIN BINDS ADENINE EITHER AS SUBSTRATE OR COFACTOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oxidoreductase; Monooxygenase; Auxin biosynthesis; Crown gall tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bonnard G., Vincent F., Otten L.; "Sequence of Agrobacterium tumefaciens biotype III auxin genes."; Plant Mol. Biol. 16:733-738(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Otten L., de Ruffray P., Submitted (FEB-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rimental princip. Sprotecia; alpha subdivision; Rhizobiaceae group; Rhizobiaceae; Rhizobium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 28.6%; Score 62; DB 1; Length 755; Local Similarity 41.7%; Pred. No. 1.9; es 15; Conservative 6; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    755 AA; 83947 MW; 9FD2B83FEA001A4D CRC64;
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P -> A (IN PTIA6NC).
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Iryptophan 2-monooxygenase (EC 1.13.12.3).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         223 DSG-----RIGFFPEDVPKPKVAVIGAGISGLVVA 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7 DSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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01-MAR-1992 (Rel. 21, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Agrobacterium vitis (Rhizobium vitis).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; K02554; AAA92550.1; -.
PIR; A04497; QQA641.
Interpro; IPR002937; Amino_oxidase.
Interpro; IPR000624; Glycosidase.
Interpre; IPR000624; Glycosidase.
Interpre; IPR000624; IND_binding.
PEm; PF01593; Amino_oxidase; 1.
Pfam; PF01593; Amino_oxidase; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=91329707; PubMed=1868204;
PLASMID=pTiA6NC;
MEDLINE=84170374; PubMed=6584906;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               r-DNA; Plasmid
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P25017;
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SO PRESENTATION OF THE PROPERTY OF THE PROPERT
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01-MAY-1991 (Rel. 18, Last sequence update)
15-JUN-2002 (Rel. 41, Last namotation update)
Genome polyprotein [Contains: Nuclear inclusion protein B (NI-B) (NIB)
(RNA-directed RNA polymerase) (EC 2.7.7.48); Coat protein (CP)]
-!- PATHWAY: FIRST STEP IN THE BLOSYNTHESIS OF AUXINS FROM TRYPTOPHAN.
-!- SIMILARITY: SIGNIFICANT HOMOLOGY TO THE ADENINE BINDING REGION OF
P-HYDROXYBENZOTHE HYDROXYLASE FROM P. FLUORESCENS. IT SEEMS THAT
THIS PROTEIN BINDS ADENINE BITHER AS SUBSTRATE OR COFACTOR.
-!- SIMILARITY: STRONG, WITH ITS COUNTERPARTS FROM A.TUMEFACIENS
PLASMIDE PITISSS, PILACHS AND PITAGNC.
-!- CAUTION: THE PLASMID PITIM4 CARRIES TWO T-REGIONS, THE TA AND
TB REGION, BOTH OF WHICH HAVE A FUNCTIONAL IAAM GENE, WITH LOW
HOMOLOGY BETWEEN THEM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR002937; Amino_oxidase.
InterPro; IPR000204; Glycosidase.
InterPro; IPR000205; NAD binding.
Pfam; PP01593; Amino_oxidase; 1.
Pfam; PF02027; RolB_RolC; 1.
Oxidoreductase; Monooxygenase; Auxin biosynthesis; Crown gall tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Viruses; ssRNA positive-strand viruses, no DNA stage; Potyviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=91016851; PubMed=2216735;
Zhou X.R., Fang R.X., Wang C.Q., Mang K.Q.;
"CDNA sequence of the 3'-coding region of PVY genome (the Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic Acids Res. 18:5554-5554(1990).
-!- ÇATALYTIC ACTIVITY: N nucleoside triphosphate = N diphosphate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              POLYPROTEIN WHICH UNDERGOES POSTTRANSLATIONAL PROTEOLYTIC PROCESSING RESULTING IN THE PRODUCTION OF AT LEAST EIGHT INDIVIDUAL PROTEINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     755 AA; 83972 MW; 6FA63E502343136F CRC64;
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es 15; Conserve
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Query Match
26.3%;
Best Local Similarity 61.1%;
Matches 11; Conservative
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354 3
378 AA;
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                                                                                                                                                                                                  RESULT 15
SLG1_YEAST
ID SLG1_YEAST
AC P54867;
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CARBOHYD
SEQUENCE
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modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (see http://www.isb-sib.ch/announce/or send an email to licenseelib-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Complete sequence analysis of the genome of the bacterium Mycoplasma
                                                                                                                                                                                                                                                                                                                              Gaps
                                                                      EMBL; X54058; CRA37993.1; -.
PIR; S11549; S11549.
InterPro; IRR001592; Poty_coat.
Pfam; PF00767; Poty_coat; 1.
Transferase; RNA-directed RNA polymerase; Coat protein; Polyprotein.
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic Acids Res. 24:4420-4449(1996).
-!- SUBCELLULAR LOCATION: Integral membrane protein (Potential).
-!- SIMILARITY: TO M.GENITALIUM MG225.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mycoplasma pneumoniae.
Bacteria, Firmicutes, Mollicutes, Mycoplasmataceae, Mycoplasma.
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STRAIN-ATCC 29342 / M129;
MEDLINE-97105895; PubMed-8948633;
Himmelreich R., Hilbert H., Plagens H., Pirkl E., Li B.-C.,
                                                                                                                                                                                                                                                                                 Match 26.3%; Score 57; DB 1; Length 327; Local Similarity 53.1%; Pred. No. 3.7; es 17; Conservative 0; Mismatches 5; Indels
                                                                                                                                                                                                    NUCLEAR INCLUSION PROTEIN B.
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                                                                                                                                                                                                                           61 327 COAT PROTEIN.
327 AÅ; 36868 MW; 8F8355E2DE6F2F18 CRC64;
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01-NOV-1997 (Rel. 35, Last sequence update)
116-OCT-2001 (Rel. 40, Last annotation update)
MPDOThetical protein MG226 homolog (F10_orf503).
MPN319 OR MPS17.
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InterPro; IPR002293; AA/rel prmeasel.
InterPro; IPR04841; Permease.
Pfam; PF00324; aa permeases; 1.
Hypothetical protein; Transmembrane; C
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468
503 AA;
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SEQUENCE
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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01-0CT-1996 (Rel. 34, Last sequence update)
01-NOV-1997 (Rel. 35, Last annotation update)
01-NOV-1997 (Rel. 35, Last annotation update)
SLG1 protein precursor.
SLG3 DR YOROOSC OR UNF378.
Saccharomyces crevisiae (Baker's yeast).
Saccharomycetales; Saccharomycetes;
Saccharomycetales; Saccharomycetes.
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SLGI PROTEIN.
N-LINKED (GLCNAC. . .) (POTENTIAL.)
N-LINKED (GLCNAC. . .) (POTENTIAL.)
EEE164F2374CCCE3 CRC64;
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SEQUENCE FROM N.A.

Sterky F., Holmberg A., Pettersson B., Uhlen M.;

Sterky F., Holmberg A., Pettersson B., Uhlen M.;

"The sequence of 30 kb fragment on the left arm of chromosom
"The sequence of 30 kb fragment con the left arm of chromosom
"The sequence of 30 kb fragment on the left arm of chromosom
"The sequence of 20 kb fragment on the left arm of chromosom
"The sequence of 10 kb fragment on the left arm of chromosom
"The sequence of 10 kb fragment of
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Length 503;
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               DB 1;
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; Pred. No. 5;
5; Mismatches
       Score 57; DB 1
Pred. No. 5.7;
3; Mismatches
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SGD; SO05534; SLG1.

InterPro; IPRO02889; WSC.

FRAM; SMO0321; WSC; 1.

Glycoprotein; Signal.

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Best Local Similarity 42.4%;
Matches 14; Conservative
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| 131 VKDNNGALIGLLVGGFVL 148
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091963 xenopus. ap 098619 xenopus lae 057394 narke japon 057394 narke japon 05923 homo sapien 073279 fugu rubrip 093279 fugu rubrip 090028 brachydanio 090028 brachydanio 090028 brachydanio 090028 brachydanio 090028 homo sapien 097917 bos taurus 097917 bos taurus 097917 agrobacteri 09173 agrobacteri 09173 agrobacteri 09163 arabiopsis

Title: Perfect score:

Sequence:

OM protein

Run on:

Scoring table:

Searched:

Minimum DB 8 Maximum DB 8

Database

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MEDINE 93236601; PubMed-8476439;
MEDINE 93236601; PubMed-8476439;
MEDINE 93236601; PubMed-8476439;
MEDINE 93236601; PubMed-8476439;
MEDINE 9325601; PubMed-8476439;
MEDINE R.B., Rosenzewaig R., Miller D.L.;
MEDINE R.B., Rosenzewaig R., Miller D.L.;
MEDINE Res. Commun. 192:96-103(1993).
MEDINE Res. Commun. 192:96-103(1993).
MESSP; PoSO67; LBA4.
INTERPRO; IPRO1255; Beta-APP.
Fram; PPO3494; Beta-APP; 1.
NON TER
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016014; Old 16014
01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 21, Last annotation update)
Beta-amyloid peptide (Fragment).
Homo sapiens (Human).
Homo sapiens (Human).
Mammalla; Metazoa, Chordata, Craniata; Vertebrata; Buteleostomi; Mammalla; Butheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.0%; Score 217; DB 4; Length 82; Best Local Similarity 100.0%; Pred. No. 1.9e-22; Matches 42; Conservative 0; Mismatches 0; Indels
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Q8UUR9
Q9UCD1
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Q9UCC8
Q8WZ99
Q9RPS4
Q8RZ41
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093279
Q9UCA9
Q8UUI8
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Q44388
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022662
Q91963
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Q919E7
Q90W28
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Q16019 homo sapien
Q16020 homo sapien
P78438 homo sapien
O93296 gallus gall
Q9pvll gallus gall
Q95kn7 macaca fasc
Q60496 cavia sp. p
Q9dgj8 gallus gall
Q9dgj7 gallus gall
Q9dui0 sus scrofa
Q11778 homo sapien
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099k32 mus musculu
098sg0 xenopus lae
P97487 mus musculu
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                                                                                                                                                                                                                                                           1 DAEFRHDSGYEVHHQKLVFF.....DVGSNKGAIIGLMVGGVVIA
                     GenCore version 5.1.4 p5_4578
Copyright (c) 1993 - 2003 Compugen_Ltd.
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Listing first 45 summaries
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Q16019
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SEQUENCE OF 32-63 FROM N.A.
MEDLINE=93035397; PubMed=1415269;
Kamino K., Orr H.T., Payami H., Wijsman E.M., Alonso M.E., Pulst S.M.,
Anderson L., O'dahl S., Nemens E., White J.A.;
"Linkage and mutational analysis of familial Alzheimer disease
kindreds for the APP gene region.";
Am. J. Hun. Genet. 51:998 -1014 (1992).
EMBL; M29270; AAA51768.1;
EMBL; M29273; AAA51768.1;
EMBL; M29273; AAA51768.1;
EMBL; M29273; AAA51768.1;
HSSP; P05067; LBA4.
                                                                                                                                                                                                                                                                                                                                                   "Alzheimer's disease amyloid peptide is encoded by two exons and shows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-NoV-1998 (TrEMBLrel. 08, Created)
01-NoV-1998 (TrEMBLrel. 08, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
Amyloid protein (Fragment).
Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE OF 19-48 FROM N.A. MEDLINES 120329; PubMed=2949367; MEDLINES 120329; PubMed=2949367; Tanzi N.E. Goorge-Hyslop Yanzi N.E., Gusella J.F., Warkins P.C., Bruns G.A., George-Hyslop Van Keuren M.L., Patterson D., Pagan S., Kurnit D.M., Neve R.L.; Mamyloid beta procein gene: cDNA, mRNA distribution, and genetic linkage near the Alzheimer locus."; Science 235:880-884(1987).
                                                                                                                                                                     Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Butaleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                               MEDLINE-89392030; PubMed=2675837;
Johnstone E.M., Chaney M.O., Moore R.E., Ward K.E., Norris F.H.,
Little S.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A. MEDDMed=9671674; MEDDINE=98337885; PubMed=9671674; Barnes N.Y., Li L., Yoshikawa K., Schwartz L.M., Oppenheim R.W. Malligan C.E.; "Increased production of amyloid precursor protein provides a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 217; DB 4; Length 82; 100.0%; Pred. No. 1.9e-22; ive 0; Mismatches 0; Indels
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                                       01-MAY-1997 (TrEMBLrel. 03, Created)
01-MAY-1997 (TrEMBLrel. 03, Last sequence update)
01-MNY-2002 (TrEMBLrel. 21, Last annotation update)
Amyloid protein (Beta-amyloid protein) (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                        similarity to soybean trypsin inhibitor.";
Biochem. Biophys. Res. Commun. 163:1248-1255(1989)
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Pfam; PF03494; Beta-APP; 1.
NON TER.
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PRELIMINARY;
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                                                                                                                                                       Homo sapiens (Human)
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                                                                                                                                                                                                                   NCBI_TaxID=9606;
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                       P78438;
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                                                                                                                                                                                                                                 SEQUENCE FROM N.A.

MEDLINE-39236601; PubMed=8476439;
Denman R.B., Rosenzcwaig R. fediler D.L.;
"A system for studying the effect(s) of familial Alzheimer disease
mutations on the processing of the beta-amyloid peptide precursor.";
Biochem. Biophys. Res. Commun. 192:96-103(1993).

EMBL; S61380; AAB26264.2;
HSSP; POS067; 1BA4.
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MEDLINE=9323661; PubMed=8476439;
MEDLINE=9323661; PubMed=8476439;
Denman R.B., Rosenzcwaig R., Miller D.L.;
A system for studying the effect(8) of familial Alzheimer disease mutations on the processing of the beta-amyloid peptide precursor.";
Biochem. Biophys. Res. Commun. 192:96-103(1993).
Biochem. Biophys. Res. Commun. 192:96-103(1993).
HSSP; P05067; 1BA4.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     82 AA; 8938 MW; F534AA50E579230A CRC64;
                01-NOV-1996 (TrEWBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
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Q16020;
01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Beta-amyloid peptide (Fragment).
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                                                                                     Beta-amyloid peptide (Fragment).
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Pfam; PF03494; Beta-APP; 1.
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Homo sapiens (Human)
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WEDLINES-9720-6426; PubMed=9116031;

WEDLINES-9720-6426; PubMed=9116031;

WEDLINES-9720-6426; PubMed=9116031;

AMEDLINES-9720-6426; PubMed=9116031;

Beck M., Mealler D., Biql V.;

"Amyloid precureor protein in Guinea pigs - complete cDNA sequence and

"Anyloid precureor protein in Guinea pigs - complete cDNA sequence and

"Anyloid precureor protein in Guinea pigs - complete cDNA sequence and

"Anyloid precureor protein in Guinea pigs - complete cDNA sequence and

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SEQUENCE PROM N.A.
TISSUE=CEREBELLUM;
MEDILINE=91273117; PubMed=1905108;
MEDILINE=91273117; PubMed=1905108;
Podlisny M.B., Tolan D.R., Selkoe D.J.;
Podlisny M.B., Tolan D.R., Selkoe D.J.;
"Homology of the amyloid beta protein precursor in monkey and human supporters a primate model for beta amyloidosis in Alzheimer's disease.";
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Hystricognathi; Caviidae; Cavia.
Amyloid b-protein precursor.
Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
Ebkaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Cercopithecidae;
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100.0%; Score 217; DB 11; Length 695;
Best Local Similarity 100.0%; Pred. No. 2.4e-21;
Matches 42; Conservative 0; Mismatches 0; Indels 0;
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Best Local Similarity 100.0%; Pred. No. 2.4e-21;
Matches 42; Conservative 0; Mismatches 0; Indels C
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01-NOV-1996 (TYEMBLrel. 01, Last sequence update)
01-JUN-2002 (TYEMBLrel. 21, Last annotation update)
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EMBL, MS9727, AAA36629. Li.
InterPro; IPR001869, A4 APP.
InterPro; IPR001255, BeTa-APP.
Fram: PP02177, AE EXTRA, 1.
Fram: PF03494; Beta-APP; 1.
PROSITE; PS00319; A4 EXTRA; UNKNOWN 1.
PROSITE; PS00320; A4 INTRA; UNKNOWN 1.
STGNAL.
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                                                                                                                                                                                           Cercopithecinae; Macaca.
NCBI_TaxID=9541;
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TISSUES ABRAIN;

Coulson B.J. Paliga K., Beyreuther K., Masters C.L.;

TOURSON B.J. Paliga K., Beyreuther K., Masters C.L.;

That the evolution of the amyloid protein precursor supergene family
tells us about tes function.";

The second of the amyloid protein precursor supergene family
tells us about tes function.";

Meurochem. Int. 0:0-0(2000).

EMBL; AF030341; AAPP.

REMBL; AF030341; AAPP.

RESPECTOR IPRO01255; BeTa-APP.

RESPECTOR IPRO01265; BeTA-APP.
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(Gallus gallus (Chicken),

Gallus gallus (Chicken),

Eukaryota; Metazpa; Chordata; Craniata; Vertebrata; Euteleostomi;

Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;

Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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0
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100.0%; Score 217; DB 13; Length 534;
Best Local Similarity 100.0%; Pred. No. 1.8e-21;
Matches 42; Conservative 0; Mismatches 0; Indels 0;
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Best Local Similarity 100.0%; Pred. No. 1.96-21;
Matches 42; Conservative 0; Mismatches 0; Indels 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NON_TER 1 -1
SEQUENCE 569 AA; 64753 MW; 0AB8BB851863A19D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NON TER 1 1 SEQUENCE 534 AA; 60597 MW; PB53ECC2E66D4C92 CRC64;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MN-2002 (TrEMBLrel. 21, Last annotation update)
Amyloid protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   436 DAEPRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMYGGVVIA 477
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Last sequence update)
Last annotation update)
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substrate for caspase-3 in dying motoneurons.";
J. Neurosci. 18-5869-5880(1998).
EMES. 1960-1989. AAC25052.1;
HSSP; P050-7; 1BA4.
InterPro; 1PR001265; BeTa-APP.
InterPro; 1PR001255; BeTa-APP.
Ffam; PP02177; A4_EXTRA; 1.
Pfam; PP03177; A4_EXTRA; 1.
PROSITE; PR003103; AMYLOIDA4.
PROSITE; PS003103; A4_EXTRA; 1.
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Q95KW7;
01-DEC-2001 (TEMBLrel. 19,
01-DEC-2001 (TEMBLrel. 19,
01-JUN-2002 (TEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
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Q95KN7
ID Q95KD
AC Q95KD
DT 01-DE
DT 01-DE
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Q9PVL1
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RESULT 9
09DGJ8
1D 09DGJ8
AC 09DG,
DT 01-M
DT 01-M
DT 01-M
DT 01-M
DT 01-M
DE Beta
OC Archa
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Amyloid precursor protein.
Sus acrofa (pig).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Cetartiodactyla, Suina, Suidae, Sus
                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 217; DB 13; Length 751; Best Local Similarity 100.0%; Pred. No. 2.7e-21; Matches 42; Conservative 0; Mismatches 0; Indels 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A Kimura A., Takahashi T.;

Kimura A., Takahashi T.;

Kimura A., Takahashi T.;

Submitted (Step-1999) to the EMBL/GenBank/DDBJ databases.

EMBL; AB032550; BAA84580.1; -.

REMBL; AB032550; BAA84580.1; -.

RICEPPO; IPR001868; A4 APP.

RICEPPO; IPR002255; Baca AAP.

RICEPPO; IPR002255; Maria EBPTI.

R Pfam; PF00147; A4 EXTRA; 1.

R Pfam; PF00147; A4 EXTRA; 1.

R PRINTS; PR00205; RANICIDEAS.

R PRINTS; PR00205; RANICE_BPTI; 1.

R PRNSTTS; PR00213; A4 EXTRA; 1.

R SWART; SM0006; A4 EXTRA; 1.

R PROSITE; PS00280; A4 EXTRA; 1.

R PROSITE; PS00280; BATIKUNITZ_1; 1.

R PROSITE; PS00280; BPTI KUNITZ_2; 1.
SMART; SM00006; A4_EXTRA, 1.

PROSITE; PS001319; A4_EXTRA, 1.

PROSITE; PS00320; A4_INTRA, 1.

PROSITE; PS00280; BPTI KUNITZ_1; 1.

PROSITE; PS0279; BPTI KUNITZ_2; 1.

Serine procease inhibitor.

SEQUENCE 751 AA, 84705 NW; E78E9413A8033D84 CRC64;
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86961 MW; SF7AlDCB2BCC583E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             653 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 694
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-WAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 DAEFRHDSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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SEQUENCE 770 AA; 86
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Q9TUIO
ID Q9TUI
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Q13778.
ID Q13778
AC Q137778
DT 01-NOI
DT 01-UND
         S KW B B KW 
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Ol-MAR-2001 (TrEMBLrel. 16, Last sequence update)

Ol-MAR-2001 (TrEMBLrel. 16, Last sequence update)

Ol-MAR-2002 (TrEMBLrel. 21, Last annotation update)

Beta-amploid precursor protein 695 isoform.

Gallus gallus (Chicken).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
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Q9D
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"Cloning of full-length chicken beta-amyloid precursor protein isoforms.";

"Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.

E MBL: AF289219; AAG00594.1;

E HSEP: P05067; 1BA4.

R HRSEP: P05067; 1BA4.

R InterPro: IPR001255; BeTa-APP.

R InterPro: IPR001255; BeTa-APP.

R InterPro: IPR001255; BeTa-APP.

R PFam; PP02177; A4 EXTRA, 1.

R PFam; PP03194; BeTa-APP; 1.

R PFam; PP03194; BeTa-APP; 1.

R PFam; PR0014; Kunitz_BPTI; 1.

R PRINTS; PR00759; BALTOFPABE.

R PRINTS; RR00759; BALTOFPABE.

R PRINTS; RR00759; BALTOFPABE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sarasa M., Rodolosse A., Sorribas V.; "Cloning of full-length chicken beta-amyloid precursor protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 217; DB 13; Length 695; Best Local Similarity 100.0%; Pred. No. 2.4e-21; Matches 42; Conservative 0; Mismatches 0; Indels 0;
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submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF289-18, AAG00593.1;
HSSP; P05067; 1BA4.
HSSP; P05067; 1BA4.
InterPro; IPR001256; Beta-APP.
Ffam; PF02177; A4_EXTRA; 1.
R PF12177; A4_EXTRA; 1.
R PRINTS; PR00303; A4_EXTRA; 1.
R PRNSTTS; PR003030; A4_EXTRA; 1.
R PROSITE; PS003129; A4_EXTRA; 1.
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                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                          PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gallus.
NCBI_TaxID=9031;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anuza; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID=10090;
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llarity 88.1%; Pred. No. 1e-18;
Conservative 4; Mismacches 1; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 91.2%; Score 198; DB 11; Length 607; Best Local Similarity 92.9%; Pred. No. 8.8e-19; Matches 39; Conservative 1; Mismatches 2; Indels C
                                                                                                                                                          NON TER 1 1
SEQUENCE 607 AA; 68391 MW; BF802214CBA7D172 CRC64;
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SEQUENCE 693 AA, 78568 MW; CAFIDF655C1AB653 CRC64;
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Van den Hurk W.H.,

Van den Hurk W.H.,

Thesis (2001), Department of Biological Sciences,
University of Nijmegen, Nijmegen, Netherlands.

EMBL, AJ298150; CA371931.j.

HSSP, P03067, 1H32.

ENTERPRO, 1H20.

InterPro, 1FR001468; A4 APP.

InterPro, 1FR001468; A4 APP.

Fram: PF03149; Beta-APP, 1.

PRINTS, PR00203; AMYLOIDA.

PROSTIT: PS00319; A4 EXTRA, 1.

PROSTIT: PS00319; A4 EXTRA, 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Beta-amyloid precursor protein A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
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hes 37; Conserv
                                                                                                                                            SEQUENCE FROM N.A.
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                                                                                                                             MEDILINE=87120328; PubMed=3810169;
MEDILINE=87120328; PubMed=3810169;
MCOldgaber L. Lerman M.I., McEride O.W., Saffiotti U., Gajdusek D.C.;
Goldgaber L. Lerman M.I., McEride O.W., Saffiotti U., Gajdusek D.C.;
Tharacterization and chromosomal localization of a cDNA encoding or construction and chromosomal localization of a cDNA encoding or construction and chromosomal localization of a cDNA encoding or construction and chromosomal localization of a cDNA encoding or construction and chromosomal localization of a cDNA encoding or construction of Alba and Chromosomal Localization of a cDNA encoding or construction or construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cricetulus griseus (Chinese hamster).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata; Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Cricetinae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 97;
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95.4%; Score 207; DB 4; Length 97

Best Local Similarity 100.0%; Pred. No. 5.59-21;

Matches 40; Conservative 0; Mismatches 0; Indels
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Sambamurti K., Pinnix I., Gandhi S.;
Sambamurti K., Pinnix I., Gandhi S.;
Sumbamurti K., Pinnix I., Gandhi S.;
BWBL; AF030413; AAB86608.1; -.
HSSP; P05667; 1BA4.
HSSP; P05667; 1BA4.
TheerPro; 1PR001255; Beta-APP.
Ffam; PF03494; Beta-APP; 1.
NON TER 19 79
SEQÜENCE 79 AA, 8538 MW; 37F2C6C3BFF3F597 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                099K32;
01-UNN-2001 (TrEMBLrel. 17, Created)
01-UNN-2001 (TrEMBLrel. 17, Last sequence update)
01-UNN-2002 (TrEMBLrel. 21, Last annotation update)
Hypothetical 68.4 kDa protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Alzheimer's amyloid beta protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 DAEFRHDSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 EFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 EFRHDSGYEVHHQKLVFFAEDVGSNKGALIGLMVGGVVIA 42
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035463
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099K32
ID 099K3;
AC 099K3;
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DE HYPOT!
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Search completed: April 21, 2003, 12:07:22 Job time : 30 secs

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Human amyloid beta
Human amyloid prot
Human amyloid-beta
Human amyloid pept
Beta/A4-amyloid pe
Human amyloid beta
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Amyloid-beta (Abet
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Amino acids 1-42 o
Human amyloid beta
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Beta-amyloid (1-43
Amyloid beta-prote
Beta amyloid prote
A-beta protein (43
Human amyloid beta
Human beta-amyloid
A-beta-binding pep
Mutant aggregating
Human tachykinin a
Synthetic amyloid
Beta-amyloid 1-42
                                                                             Beta-amyloid 1-42
Amyloid-beta pepti
Human peptide anti
Human Alzheimer-be
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Amyloid beta-prote
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Neuronal death inh
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Region of pre-AP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transgenic mice as models for studying Alzheimer's disease proteins - contg. cells with promoter and beta-amyloid precursor protein deoxyribonucleic acid, useful for testing anti-alzheimer's drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transgenic mice; Alzheimer's disease; diagnosis; beta-amaloid precursor; plaque core protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence of A99 (beta-amyloid core domain).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ALIGNMENTS
                           AAW29093
AAW99268
AAW81474
AAY96956
AAY96956
AAB8622
AAB81623
AAB91779
AAB91812
AAB48999
AAB48999
AAB48999
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ABB83306
AAU96896
ABB81321
AAE21438
ABB76029
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AAU80961
AAG68314
AAM51864
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AAR64165
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     (CALB-) CALIF BIOTECHN INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1992-024426/03.
    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JUN-1990;
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     AAR20330;
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AAR20330
ID AAR2
       Sequence of A99 (b
Beta-amyloid (1-42
Beta/A4-amyloid pe
Alzheimer amyloid
Beta A4 peptide.
Neurotoxic beta-am
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Full length beta-a
Human beta amyloid
Human amyloidogeni
                                                                                         April 21, 2003, 11:57:04 ; Search time 36 Seconds (without alignments) 155.459 Million cell updates/sec
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11. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1880.DAT:*

22. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1881.DAT:*

33. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1881.DAT:*

34. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1881.DAT:*

35. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1881.DAT:*

36. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1881.DAT:*

37. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1881.DAT:*

38. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1881.DAT:*

39. ($1D52/gcgdata/geneseq/geneseqp-emb1/AA1881.DAT:*

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                                                                                                                                                                                                                                                                    908470
                 GenCore version 5.1.4 p5_4578
Copyright (c) 1993 - 2003 Compugen_Ltd.
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Maximum Match 100%
Listing first 45 summaries
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AAR95248
AAR94591
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AAW44507
AAW47230
AAW4729891
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1 DAEFRHDSGYEVHHOKLVFF.
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WPI; 1996-268332/27.
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                                                                                                                                                                                                                                                                                                           Homo sapiens
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                           The inventors specifically claim transgenic mice contg. DNA encoding A42 (beta-amyloid precursor protein) (AAR2030), A99 (beta-amyloid carboxy tail) (AAR20329), A695 (beta-amyloid precursor protein), A751 (precursor plus inhibitor) or A41 (protease inhibitor) (AAR20328). Human fibroblast cDNA clone lambdaAPCP10814 was deposited at ATCC on July 1, 1987 and has accession No. 40347. The promoter is pref. the NSE promoter with the A751 or the A695 sequence.
                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibodies which recognise specific subfragments of the beta-amyloid protein are claimed. Specifically, the antibodies (which are pref. monoclonal) recognise residues 1-16 and/or 1-28 from the N-terminal portion of beta-amyloid or they recognise residues 25-35 or 35-43 from the C-terminal portion. The antibodies are useful for assaying beta-amyloid and its derivatives for diagnosis of Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amyloid precursor protein, APP, Alzheimer's disease, beta-amyloid, anti-beta-amyloid antibody, diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antibodies recognising specific parts of beta-amyloid - can be used for diagnosis of diseases implicating beta-amyloid, such as Alzheimer's disease
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0
                                                                                                                                                                        100.0%; Score 217; DB 13; Length 42; 100.0%; Pred. No. 8.2e-25; tive 0; Mismatches 0; Indels
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   Disclosure, Fig 3; 98pp; English.
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93JP-0019035.
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                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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42; Conservative
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Best Local Similarity
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05-FEB-1993;
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To investigate the effects of beta-amyloid peptide (BAP) on tissue plasmingen activator (t-PA) 3 synthetic peptides were used. Con peptide contained 42 amino acids and corresp. to the full one peptide contained 42 amino acids and corresp. to the full length BAP (AAR95248). The other 2 peptides (AAR95249 and 50) contained the 28 Arceminal residues of the BAP found in Alzhaimer's disease and hereditary cerebral haemorrhage with amyloidosis-Dutch type and hereditary cerebral haemorrhage with amyloidosis-Dutch type (HCHWA-D), respectively. In an assay to determine the effect of the peptides on t-PA activation, each peptide (AAR95248, 49 and 50) gave 1st order rate constant of activation (k(app)) values of 31.4, 13.9 and 41.5, respectively, compared to 1.7 and 7.8 for nill and fibrinogen controls. The results demonstrate that the BAP are able to stimulate t-PA activity in vitro, which is significant in that it provides a means for investigating and controlling the pathogenesis of Alzheimer's disease, HCHWA-D and cerebral amyloid angiopathy related cerebral haemorrhage.
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                                                                                                                                                                                                                                                                                                                                              Beta/A4-amyloid peptide; tissue plasminogen activator;
Alzheimer's disease; stimulation; investigation; pathogenesis;
hereditary cerebral haemorrhage with amyloidosis-Dutch type;
control; cerebral amyloid anglopathy; cerebral; haemorrhage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Use of agents which bind beta-amyloid peptide - for diagnosis, prevention and treatment of vascular damage caused by amyloid deposits, partic. in haemorrhaging and Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 217; DB 17;
100.0%; Pred, No. 8.2e-25;
iive 0; Mismatches 0;
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AAR95248 standard; peptide; 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94US-0347144.
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                                                                                                                                                                              (first entry)
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Best Local Similarity 100.
Matches 42; Conservative
                                                                                                                                                                                                                                                            Beta/A4-amyloid peptide.
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Gaps

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Indels

us-09-580-018-42.rag

Synthetic.

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This sequence represents the beta A4 peptide. The degree of aggregation of this peptide is determined using the method of the invention. The beta h4 peptide is present in the brain of Alzheimer's disease patients, but not in the brain of horalmer's disease patients, but clumps or aggregates in the brain of Alzheimer's disease patients, where it may be responsible for the destruction of normal brain cells. Once the clumps or aggregates form, the formulation is almost irreversible. The method of the invention comprises reacting this sequence with a binding reagent capable of binding to it only in its non-aggregated state, to form an amount of a beta A4 peptide-bound reagent and an amount of percent of the beta A4 peptide, binding reagent complex is then measured. Compounds which inhibit aggregation of beta A4 peptide are potentially useful for treatment of Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beta-amyloid peptide; beta-AP; neuropeptide; neurotoxin; calcium influx; aggregate; Alzheimers disease; decoy; treatment.
                                                                                                                                                                                                                                                                          Determn. of the degree of betaA4 peptide aggragation using binding agent - used to screen cpds, for possible use in Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 217; DB 18; Length 42; 100.0%; Pred. No. 8.2e-25;
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                                                                                                                                                                                              Sahasrabudhe SR
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                                                                                                                                                      (HMRI ) HOECHST MARION ROUSSEL INC.
                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 10; 18pp; English.
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                                                                                                                                                                                              Paul JW, Riedel NG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW64507 standard; peptide; 42
                                                                           96WO-US12034.
                                                                                                                 95US-0515606.
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97US-0035847
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                                                                                                                                                                                                                                  WPI; 1997-165447/15.
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les 42; Conserv
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WO9707403-A1.
                                                                           23-JUL-1996;
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                                                                                                                 16-AUG-1995;
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                                                                                                                                                                                                                                                                                                                treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                    Beta-amyloid; Alzheimer's disease; cholinesterase; lipase; ebelactone-A; serine protease; para-amidinophenylmethanesulphonyl fluoride; inhibition; complex formation; alpha(1)-antichymotrypsin; Down's diseases; ageing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This is the sequence of a fragment of the beta-amyloid protein associated with Alzahaimer's disease. The protein contains esterase (cholinesterase and lipase) activities based on active site similarities with serine proteases (see AAR94552-96). The esterase activity of the beta-amyloid protein is inhibited by the cpds. of the invention i.e. belactone A or para-amidinophenylmethanesulphonyl fluoride. Inhibition of these activities prevent complex formation between the beta-amyloid protein and alpha(1)-antichymotrypsin, thus can be used to treat, study or diagnose Alzheimer's or Down's diseases or normal ageing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inhibiting enzymatic activity of Alzheimer amyloid beta-protein using p-amidino:phenyl:methanesulphonyl fluoride or ebelactone A, for treatment, study and diagnosis of Alzheimer's disease, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beta A4 peptide; alzheimer's dísease; peptide aggregation; brain;
therapy; inhibitor.
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                                                                                                            Alzheimer amyloid beta-protein actíve site sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Score 217; DB 17;
; Pred. No. 8.2e-25;
0; Mismatches 0;
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AAR94591 standard; peptide; 42 AA.
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ilarity 100.0%;
Conservative 0;
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90US-0572671,
92US-0819361,
93WO-US00325,
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                                                                         21-AUG-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HARD ) HARVARD COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Potter H;
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Best Local Similarity
Matches 42; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           42 AA;
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13-JAN-1992;
13-JAN-1993;
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AAW12828;

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Length 42;

DB 19;

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Beta-amyloid peptide; BAP; extracellular BAP plaque;
cerebrovascular deposit; Alzheimers disease; Downs syndrome;
amyloid precursor protein; APP; secretase; BAP aggregation;
                                                                                          1 DAEFRHDSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
                                                                                                        1 DAEFRHDSGYEVHORLVFFAEDVGSNKGAIIGLMVGGVVIA 42
                                     1100.0%; Score 217; DB 19;
1100.0%; Pred. No. 8.2e-25;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                           Full length beta-amyloid peptide (BAP).
                                                                                                                                                                                AAW42989 standard; peptide; 42
                                                                                                                                                                                                                                                                                                            amyloid precursor protein, API
abnormal proteolytic cleavage
                                                                                                                                                                                                                                  (first entry)
                                 Ouery Match
Best Local Similarity 100.0
Marches 42; Conservative
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               42 AA;
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                                                                                                                                                                                                                                 01-MAY-1998
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01-MAY-1992;
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                                                                                                                                                                                                         AAW42989;
               Sequence
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                                                                                                   AAW64488-W64517 are decoy peptides that bind to a neurotoxic beta-amyloid peptide (beta-AP) and reduces the ability of beta-AP's to form aggregates that increase calcium influx into neuronal cells. Such peptides can be used in the treatment of diseases associated with neurocroxic aggregates of beta-AP specifically Alzheimer's disease. The peptides are administered at 0.001-1000 (especially 0.2-20) mg/kg, by injection and orally, or from slow-release implants.
                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New in vitro screening assay for Alzheimer's disease drugs -comprises assessing binding of labelled beta-amyloid peptide to silk sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence was used in the development of a novel in vitro screening assay for agents capable of affecting the deposition of beta-amyloid peptide (BAP) on tissue. The method comprises contacting a silk sample with labelled BAP, optionally in the presence of a test agent, detecting the amount of label bound to the silk and assessing the effect of the agent on the deposition of BAP. Agents that inhibit binding of BAP to silk are potentially useful for treating amyloidosis diseases, especially Alzheimer's disease.
                                                                                                                                                                                                                                                            6
                        Inhibition of aggregation of, e.g. beta-amyloid peptide - by administering decoy peptide or other calcium-influx inhibitor, useful for, e.g. treating Alzheimer's disease
                                                                                                                                                                                                                                 100.0%; Score 217; DB 19; Length 42; 100.0%; Pred. No. 8.2e-25; ive 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                  1 DAEFRHDSGYEVHHQKLVFFRAEDVGSNKGAIIGLMVGGVVIA 42
                                                                                                                                                                                                                                                                                     1 DAEFRHDSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Screening assay, beta-amyloid peptide, treatment, amyloidosis disease, Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beta-amyloid peptide residues 1-42
                                                                            Example 8; Page 46; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                         AAW47230 standard; peptide; 42
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91US-0744767.
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                                                                                                                                                                                                                                             Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Maggio JE, Mantyh PW;
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WPI; 1998-398795/34.
                                                                                                                                                                                                         42 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-SEP-1994;
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                                                                                                                                                                                                          Sequence
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amyloid precursor protein fusion polypeptides - comprising APP fragment and marker, useful for research and drug screening
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100.0%; Score 217; DB 19; Length 42;
Best Local Similarity 100.0%; Pred. No. 8.2e-25;
Matches 42; Conservative 0; Mismatches 0; Indels 0
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                                                                                                                                                                                                                     (AMCY ) AMERICAN CYANAMID CO.
                                                                                                                                92US-0877675.
                                                                                         93US-0123659.
95US-0464248.
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This invention describes a novel method for inhibiting amyloid fibrillogenesis which comprises contacting tissue with a composition comprising an amyloidogenic peptide, beta-amyloid, that has been blocked at an end terminal or a side chain, by conjugation to polyethylene glycol, by conjugation to a second compound and a pharmaceutically acceptable buffer, solvent or diluent. The methods are used to inhibit amyloid plaque formation in humans suffering from amyloidosis, Alzheimer's disease or Down's Syndrome. This sequence represents a fragment of the beta-amyloid peptide described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Amyloid protein; beta-A4 peptide; aggregation; screening; inhibition; therapeutic drug; brain; Alzheimer's disease.
        Amyloidogenic; beta-amyloid; A-beta peptide; human; inhibitor;
fibrillogenesis; amyloid plaque; amyloidosis; Alzheimer's disease;
Down's Syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                               Inhibiting amyloid plaque formation in humans suffering from amyloidosis, Alzheimer's disease or Down's Syndrome -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 217; DB 20;
100.0%; Pred. No. 8.2e-25;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                    Burkoth TS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 21; Page 140; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY25137 standard; peptide; 42 AA
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Best Local Similarity 100.v
                                                                                                                                                                                                                                                                                                                                  Meredith SC,
                                                                                                                                                                                                                                                                                           (ARCH-) ARCH DEV CORP
                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-561326/47.
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                                                                                                                                  WO9941279-A2
                                                                                                                                                                                                                 12-FEB-1999;
                                                                                               Homo sapiens
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                                                                                                                                                                          19-AUG-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention.
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                                                                                                                                                                                                                                                                                                                                    Lynn DG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a cell line which produces beta amyloid precursor protein (APP) and expresses alpha secretase activity but expresses beta secretase activity only under an external stimulus. Also described is a cloning method for DNA encoding beta secretase, comprising: (1) inserting a DNA library into the cell line, expressing the inserted DNA, and selecting cells expressing beta secretase then isolating the beta secretase DNA from them, or (2) isolating nucleic acid from the cell line with or without external stimulation and performing subtractive cloning to identify DNA expressed only under stimulation. Products from the present invention may be used in the investigation of neurological disorders such as Alzheimer's disease and Downs syndrome and in particular the association of mutations of the bar are all present sequence represents a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A cell line which produces beta amyloid precursor protein, used in the investigation of neurological disorders such as Alzheimer's disease -
                                                                                                                                                                                                                                                    Human; beta amyloid precursor protein; APP; beta secretase inhibition; alpha secretase; neurological disorder; Alzheimer's disease; Downs syndrome; mutation.
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DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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                                                                                                                                                                                                               Human beta amyloid precursor protein peptide.
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                                                                                             AAY49691 standard; peptide; 42 AA
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                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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Gaps

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Length 42; Indels

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                                                                                                                                                                                               This invention describes a novel method for the determination of the degree of aggregation of an amyloid beta A4 peptide (I) in solution. Determination comprises: (a) incubating a sample of unaggregated (I) with Coomassie Brilliant Blue G 250 dye (II) which only binds to unaggregated (I); (b) measuring the amount of (II) bound to (I) to obtain a value (i); (c) repeating steps (a) and (b) with a second sample at a different time to obtain a second value (ii); and (d) eterminate the difference between (I) and (II) which is inversely related to the degree of aggregation of (I). This method may be applied to a screen for compounds that inhibit aggregation of (I). These inhibit toss may be used as therapeutic drugs to inhibit the formation of these aggregates in the brains of patients suffering from Alzheimer's disease.
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                                                                                        Determination of degree of aggregation of a peptide, useful for identifying therapeutic drugs for treating Alzheimer's disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11arity 100.0%; Score 217; DB 20; Length 42; 11arity 100.0%; Pred. No. 8.2e-25; Conservative 0; Mismatches 0; Indels
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Sahasrabudhe SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human beta-amyloid precursor core protein A42.
                                                                                                                                                         Claim 1; Column 7-8; 8pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY08607 standard; Protein; 42 AA
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Riedel NG,
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90US-0538857.
91US-0716725.
95US-0422333.
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                                             WPI; 1999-403957/34.
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Matches 42; Conserv
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Goyal S,
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This invention describes novel transgenic mice expressing proteins related to the pathology of Alzheimer's disease and which provide models for studying potentially therapeutic compounds. The transgenic mice contain a DNA sequence encoding a beta-amyloid precursor protein (APP) and a nerve tissue specific promoter operably linked to the beta-APP allowing its expression to form beta-amyloid protein deposits in the animal's brain. The transgenic mouse is useful for elucidating the molecular mechanisms involved in the synthesis of and, more importantly, inhibiting the synthesis and deposition of beta-amyloid proteins (most importantly in the brain where plaque formation is associated with Alzheimer's disease) by inhibiting production and/or increasing cleavage after production. The transgenic animals provide useful models for studying the in vivo relationships of the proteins to each other and to other compounds being tested for their usefulness in treating Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New conjugates are disclosed which are of formula A-Z, in which: A is fill a cyclosporin A analogue described in AAW29087 or (2) an FK506 binding peptide inhibitor; and Z is a polypeptide comprising 5 or more contiguous residues of A-beta peptide. The compounds are novel chantcal inducers of dimerization which are non-immunosuppressive and which are inhibitors of A-beta peptide aggregation and deposition in amyloid plaques. The adverse consequences of amyloid plaque formation can be prevented or ameliorated by sequestering the A-beta peptide in monomeric form with a conjugate which links the A-beta to cyclophilin or FKBP, therefore providing a mechanism to minimize the amount of free A-beta
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treatment of neurological diseases e.g. Alzheimer's disease,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 217; DB 20; Length 42; 100.0%; Pred. No. 8.2e-25; tive 0; Mismatches 0; Indels (
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                                                                                                                                                                                                                                                                                                                                                                                                          42 AA;
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DAEFRHDSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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                                                                                                                                                                                   27-JUL-1991;
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                                                                                                                Homo sapiens
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                                                                                                                                      US5876948-A.
                                                                                                                                                                                                                                                                 Yankner BA;
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  AAW92726;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence corresponds to a mutant aggregating amyloid-beta peptide which can be covalently labelled with a fluorescent group. The detection or monitoring of an amyloid aggregate in a sample can be used to diagnose or detect a predisposition to Alzheimer's disease. The screening assays can be used to identify compounds for the treatment or amelioration of Alzheimer's disease or its symptoms. The fluorescent derivatives of the amyloid-beta peptide are also useful for exploring other aspects of amyloid structure.
available for fibril formation and deposition. The compounds can be used for the treatment of Alzheimer's disease, multiple sclerosis and amyotrophic lateral sclerosis.
                                                                                          Gaps
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                                                                                                                                                                                                                                                                                      Aggregation; amyloid-beta peptide; fluorescent group; detection; diagnosis; Alzheimer's disease.
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                                                                   20; Length 42;
                                                                                        Indels
                                                                                                                           1 DAEFRHDSGYEVHHQKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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                                                                                                             DAEFRHDSGYEVHHOKLVFFAEDVGSNKGAIIGLMVGGVVIA 42
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                                                                 100.0%; Score 217; DB 20;
.larity 100.0%; Pred. No. 8.2e-25;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New fluorescent labeled amyloid A-beta peptides
                                                                                                                                                                                                                                                               Mutant aggregating amyloid-beta peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 21; 50pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW92726 standard; peptide; 42
                                                                                                                                                                                           AAW99585 standard; peptide; 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Glabe C;
                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US16809
                                                                                                                                                                                                                                         (first entry)
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                                                                            Local Similarity
les 42; Conserv
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nes 42; Conserv
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                                             42 AA;
                                                                                                                                                                                                                                                                                                                       Homo sapiens.
Synthetic.
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This invention describes a method for screening compounds for inhibiting a neurotoxin. The method involves incubating tachykinin agonists with neuronal cells and a beta-amyloid peptide neurotoxin. The methods can be used for identifying compounds for treating diseases characterised by an undesirable build up of beta-amyloid protein, e.g. Alzhaimer's disease, Down's syndrome, and the syndromes of hereditary cerebral haemorrhage with amyloidosis and non-inherited congophilic angiopathy with cerebral haemorrhage. AMW92655-W92731 are tachykinin agonists derived from human beta-amyloid peptide fragments.
                                                                                                                            Alzheimer's disease; Down's syndröme; amyloidosis; human;
hereditary cerebral haemorrhage; non-inherited congophilic angiopathy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Screening for neurotoxin inhibitors - by testing compounds for their effect on beta-amyloid peptide neurotoxic effect on neuronal cells
                                                                              Tachykinin agonist; beta-amyloid; inhibition; neurotoxin; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
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Human tachykinin agonist beta-amyloid peptide fragment #72.
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nes 42; Conserv
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Asp2 recognition s Human beta amyloid Human wild-type AP Synthetic peptide Human APP derived Human APP derived Sythetic peptide f Peptide #1 used as Beta-secretase spe Human APP beta-sec

AAU07227 AAE02606 AAB66574 AAB46208

AAU06628

Beta-APP alpha-sec Beta-APP alpha-sec Beta-APP alpha-sec

AAW70868 AAR24262 AAR24261 AAY69703 AAY69704 AAY69708 AAE10654

Human wild-type AP Human amyloid prec

Human APP derived Human APP derived Beta-amyloid pepti

Human amyloidin pr

Beta-amyloid prote Human NSE promoter Beta-APP alpha-sec Human APP-Sw mutan Ruman ASP-2 beta-s

Human amyloid beta Beta-secretase pep

Immunogen for rais

fmmunogen for rais

Human beta-amyloid

Peptide product of

AAB46216
AAB6136
AAB6136
ABA878615
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AAR06606
AAR066067
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Beta-amyloid pepti Human APP A-beta 1 Beta-secretase sub Human Aspartyl pro

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Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human APP derived immunogenic peptide #8.
                                                                                                                                                                                                                                                                                                                                                                                                              AAB46212 standard; peptide; 10
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 04-APR-2001 (first entry)
 (NEUR-) NEURALAB LID.
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WO200072880-A2
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  Human APP A-beta p
Synthetic peptide
Beta-amyloid pepti
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Human APP derived
Human APP derived
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Human APP derived
Human APP derived
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                                                                  April 21, 2003, 12:06:55; Search time 28.3333 Seconds (without alignments) 47.030 Million cell updates/sec
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          GenCore version
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AAB46213
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                                 This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (Fc receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta: or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient eg. active and passive immunication. The methods are useful for prophylactic and therapeutic treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beta amyloid is synthesised as part of a larger protein referred to as amyloid precursor protein (APP), which has a number of isoforms in humans, including APP65s and APP70. The anino terminal of beta amyloid is generated by cleavage of a peptide bond of APP which in APP65s lies between Met596 and Asp597. APP forms a complex with Go, a GTP-binding protein (or "G protein") in brain. Go is made of one alpha subunit and one Beta-agamma subunit. Two isoforms of Go, known as Gol (or GoA) and Go2 (or GoB) have been identified; they have slight AA differences in their alpha subunits. The cDNA sequence and
                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying cpds. useful for treating or preventing Alzheimer's disease - by determining whether it interferes with the association of the couplone portion of amyloid precursor protein to G polypeptide
                                                                                                                                                                                                                                                                                    ;
                                                                                                                                                                                                                                                         Length 10;
                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; isoform APP 695; beta amyloid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human amyloid precursor protein APP695 residues 597-606.
                                                                                                                                                                                                                                                       100.0%; Score 10; DB 22; 100.0%; Pred. No. 2.3e-05;
                                                                                                                                                                                                                                                                                    Mismatches
            Disclosure; Figure 19; 143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                          AAR58928 standard; peptide; 10 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                    Alzheimer's disease.
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Alzheimer's disease
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les 10; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nishimoto I;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                              Gaps
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deduced AA sequence of the alpha subunits in each of Gol and Go2 shown in AAQ69002/R58914 and AAQ69004/R58924 respectively. The cytoplasmic App695 sequence His657-Lys676 (AAR58913) possesses a specific Go-activating function, and is necessary for complex formation of this APP with Go. AAR58928 is another peptide of APP695 which corresp. to AAS 597-606.
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100.0%; Pre
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Best Local Similarity 100.0%; P.
Matches 9; Conservative 0;
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                                                                                                                                                                                                                                                                             Local Similarity 100.
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Matches
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This invention relates to a novel method for modulating activity of beta-site amyloid precursor protein (APP)-cleaving enzyme (BACE)

Secretase/Sheddase. Cleavage of BACE by this enzyme results in the secretase/Sheddase. Cleavage of BACE by this enzyme results in the applied peptide Abeta which enhances the production of the amyloidogenic peptide Abeta which has been shown to be involved in the actiology of Alzehmer's disease. Inhibition of BACE secretase can be actiology of Alzehmer's disease. Inhibition of BACE secretase and be activated by administration of an antisease nucleotide molecule capable of hybridising with BACE mRNA, by using a ribozyme that targets and degrades BACE secretase mRNA, with a peptide that can interfere with confine the enzyme with BACE or using an antibody or antagonist that can function as an inhibitor of BACE secretase activation. The methods of the invention modulate the activity of BACE secretase/sheddase by corprelin, especially Alzheimer's disease. The invention of Abeta apended for identification of an agent that can alter the ability of BACE secretase to associate with and process a known substrate, this method can be used for high throughput screening of candidate molecules. The invention also comprises a method for determining whether an individual is at risk of developing a neurodegenerative disorder characterised is a process. The the contrastion of an anterior of an activation of an agent that can alter the ability of BACE can be used for high throughput screening of candidate molecules. The that the can alter the additional and an activation of an anterior of an enrodegenerative disorder characterised.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modulating activity of beta-site amyloid precursor protein-cleaving enzyme secretase/sheddase for treatment of neurodegenerative disorder characterised by generation of Abeta protein, by preventing cleavage of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            is at risk of developing a neurodegenerative disorder characterised by the generation of Abeta protein by measuring the levels of BACE C terminal cleavage products in a sample or tissue where an increase in cleavage products indicates a person at risk. The present sequence represents the N terminal of a beta amyloid protein of the
                                                                       Alzheimer's disease; beta amyloid precursor protein; mouse; BACE; beta-site APP cleaving enzyme; nootropic; neuroprotective; beta-site amyloid precursor protein (APP)-cleaving enzyme; APP; BACE secretase/sheddase; neurodegenerative disorder.
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100.0%; Pred. No. 7.8e+05;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                     (RECL-) INST RECH CLINIQUES MONTREAL.
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ID AAB46210 standard; peptide; 10 AA.
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100.08; Fr
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                                                                                                                                                                                                                                                                                                                                  01-AUG-2000; 2000CA-2313828.
                                    N terminus of beta amyloid.
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                                                                                                                                                                                                                                                                                                                                                                                                              Chretien M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-280632/32.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention.
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                                                                                                                                                                                                                                                                                                             Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yednock T;
                                                                                                                                                                                                                                                                          Human APP derived immunogenic peptide #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Figure 19; 143pp; English.
                                                                                                                                                       AAB46213 standard; peptide; 10 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU78518 standard; Peptide; 8 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     methods are useful for Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bard F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-032104/04.
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                                         1 MDAEFRHDS 9
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Best Local S
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AC AAU78
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This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient. (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                                                                                                                                                                                           Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 22; L 0.0048;
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                                                                                                                                                                                                          Yednock T;
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100.0%; Pred. No. 0.0
:ive 0; Mismatches
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                                                                                                                                                                                                          Vasquez NJ,
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                                                                             26-MAY-2000; 2000WO-US14810.
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Best Local Similarity
Matches 8; Conserva
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                                                                                                                                                                                                                                                                                                                                    specific antibody
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WO200072880-A2
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                                                                                                                        28-MAY-1999;
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                                        07-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Alzheimer's disease.
                                                                                                                 Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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100.0%; Pred. No. 0.0048;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vasquez NJ, Yednock T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human APP derived immunogenic peptide #10.
                                                                             Human APP derived immunogenic peptide #6.
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Best Local Similarity 100.00
                                    (first entry)
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                                    04-APR-2001
                                                                                                                                                                                                       Homo sapiens.
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AAB46214;

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Length 10; 0; Indels

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Alzheimer's disease (AD). In particular, the invention provides a novel mutation (T141) identified in the amyloid precursor protein (APP), APP14, which leads to a very aggressive form of AD. The mutation involves the 43rd codon of the amyloid Beta peptide (A-Beta) corresponding to the putative gamma 42.secretase cleavage site. The corresponding to the putative gamma 42.secretase cleavage site. The novel mutation alters both A-Beta40 and A-Beta42 secretion elevating the A-Beta42A-Beta40 ratio by 10-fold in vitro. Furthermore, the main amyloid blaque pathology in brains of these patients is of the diffuse or pre-amyloid type composed primarily of N-truncated A-Beta42. Dense-cored plaques although not absent, were significantly reduced. Also, the instance, in vessels as cerebral amyloid anglopathy (CAA) or senile instance, in vessels as cerebral amyloid anglopathy (CAA) or senile indicate that deposition of N-truncated A-Beta42 in one of the earliest of inciting AD either through the well-established 'amyloid cascade' or of inciting AD either through the well-established 'amyloid cascade' or paying a yet unknown mechanism(s). This sequence represents a synthetic peptide of A-Beta42 by immunising mice with the synthetic peptide.

A-Beta42 by immunising mice with the synthetic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a peptide derived from beta-amyloid precursor protein (beta-APP, see AMY0885). The peptide is a beta-amyloid peptide and is used to produce a monoclonal antibody designated antisenilin N1/7. The specification describes a method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Preventing or inhibiting progression of Alzheimer's Disease comprises use of recombinant DNA encoding an antibody specific for the N- or C-terminus of an amyloid-beta peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beta-amyloid precursor protein; beta-APP; beta-amyloid peptide; antibody; amyloid deposit; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  70.0%; Score 7; DB 23; Length 7; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beta-amyloid peptide to create a monoclonal antibody.
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Best Local Similarity 100.0
From 7; Conservative
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LTD.
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(MIND-) MINDSET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Db
                                                                                                                                                                                         This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytcsis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta; or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The methods are useful for prophylactic and therapeutic treatment of Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Neurodegenerative disorder; Alzheimer's disease; AD; T7141; APP; APP714; amyloid precursor protein; amyloid Beta peptide; A-Beta; A-Beta40; brain; A-Beta42; plaque pathology; pre-amyloid; cerebral amyloid angiopathy; honse-cored plaque; CAA; senile plaque core; amyloid cascade; murine; mouse; DAEFRHD; monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             precursor reducing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to the field of the neurodegenerative disorder of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                        Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic peptide of A-Beta residues 1-7 (DAEFRHD).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                70.0%; Score 7; DB 22; L¢ 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kumar Singh S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 21; 42pp; English.
                                                                                                                                                      Claim 59; Page 119; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA014421 standard; Peptide; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUL-2001; 2001WO-EP07830.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  De Jonghe C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-154742/20.
                                                                                                        specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 AA;
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1 DAEFRHD 7
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Synthetic.
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Alzheimer's disease.

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               method comprises administering a composition comprising a recombinant DNA molecule containing a gene encoding a recombinant antibody end specific for the N-terminus or the C-terminus of an amyloid-beta peptide, operably linked to a promoter which is expressed in the central nervous system. The recombinant antibody molecules prevent the accumulation of beta-amyloid peptides in the extracellular space, interstitial fluid and cerebrosphal fluid and the aggregation of such peptides into amyloid deposits in the brain. They also inhibit the progression of Alzheimer's disease by inhibiting the interaction of beta-amyloid peptides mediating Alzheimer's disease induced neurotoxicity and inhibiting the Alzheimer's disease induced complement activation and cytokine release involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                        Gaps
  progression of Alzheimer's disease. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody
                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                Length 8;
                                                                                                                                                                                                                                                                                                              DB 19; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yednock T;
                                                                                                                                                                                                                                                                                                            70.0%; Score 7; DB 1
100.0%; Pred. No. 7.8
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human APP derived immunogenic peptide #5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB46209 standard; peptide; 10 AA.
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or inhibition of
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                                                                                                                                                                                                                                                                                                                                Local Similarity 100.
nes 7; Conservative
                                                                                                                                                                                                                                  the inflammatory process
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                                                                                                                                                                                                                                                                          8 AA;
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                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient; (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (Fc receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta; or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                     Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody
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                                                     Length 10;
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                                                     DB 22;
. 0.069;
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100.0%; Pred. No. 0.069;
iive 0; Mismatches
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                                                     Score 7; DB 2:
Pred. No. 0.0
0; Mismatches
                                                                                                                                                                                                                                                                                               Human APP derived immunogenic peptide #11.
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                                                                                                                                                                                                            AAB46215 standard; peptide; 10 AA.
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                                           70.0%; 5c.
100.0%; Pre
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                                                                                  Conservative
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Matches 7; Conservative
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                                        Schenk DB, Bard F,
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                         10 AA;
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EFRHDSG 7
                                                                                                            1 MDAEFRH 7
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                           Sequence
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(first entry)
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                                                                                             Key
Modified-site
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5 MDAEFR 10
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Modified-site
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09-NOV-1992
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                                                                                                                                                            30-APR-1992.
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                                                                         Synthetic.
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  δλ
                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents a peptide derived from beta-amyloid precursor protein (beta-APP, see AAW70863). The peptide is a breamyloid peptide and is used to produce a monoclonal antibody. The specification describes a method for prevention or inhibition of progression of Alzheimer's disease. The method comprises administering a composition comprising a recombinant DNA molecule containing a gene encoding a recombinant antibody end-specific for the N-terminus or the certain nervous system. The recombinant which is expressed in the central nervous system. The recombinant antibody molecules prevent the accumulation of beta-amyloid peptides in the aggregation of such peptides into amyloid deposits in the brain. The aggregation of such peptides into amyloid deposits in the brain. The aggregation of beta-amyloid peptides mediating Alzheimer's disease induced neurocoxicity and inhibiting the Alzheimer's disease induced complement activation and cytokine release involved in the inflammatory process.
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                                                                                                                                                                                                                                                                                                                                           Preventing or inhibiting progression of Alzheimer's Disease - comprises use of recombinant DNA encoding an antibody specific for the N- or C-terminus of an amyloid-beta peptide
                                                                                                                 Beta-amyloid precursor protein; beta-APP; beta-amyloid peptide; antibody; amyloid deposit; Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 6; DB 19; Length 6;
Pred. No. 7.8e+05;
0; Mismatches 0; Indels
                                                                                              Beta-amyloid peptide to create a monoclonal antibody.
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                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 47; 58pp; English.
                                AAW70868 standard; peptide; 6 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60.0%; Scallarity 100.0%; Pi
Conservative 0;
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                                                                                                                                                                                                                                                97US-0041850
                                                                          (first entry)
                                                                                                                                                                                                                                                                    (MCIN/) MCINNIS P A. (MIND-) MINDSET LTD.
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                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                  WO9844955-A1.
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                                                                         04-FEB-1999
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1 DAEFRH
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                                                                                                                                                   Synthetic
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                                                     AAW70868;
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Matches
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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This synthetic substrate is based on the sequence of residues 592 to 601 of the 695 amino acid APP. Lys595 has been replaced by Gln. Amyloidin cleaves the Metr-Asp bond of this peptide but does not cleave any of the other bonds.

See AAR24260-3, AAR24266-7 and AAO24875-024887.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seubert PA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alzheimer's disease; beta amyloid precursor protein; APP; zinc; metalloprotease; hAP; protease inhibitor; APP592-601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                           Alzheimer's disease; beta amyloid precursor protein; APP; zinc; metalloprotease; hAP; protease inhibitor; APP592-601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human amyloidin protease - used for cleaving Met-Asp bond in amyloid-like substrate for identifying protease inhibitors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               60.0%; Score 6; DB 13; Length 10;
100.0%; Pred. No. 0.99;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dovey HF, Johnstone EM, Little SP, McConlogue L, Sinha S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human amyloidin protease substrate sequence #1.
Human amyloidin protease substrate sequence #2.
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                                                                                                                                                                                                                                                                                 /note= "Acetylated-Ser"
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                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Page 29; 62pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ATHE-) ATHENA NEUROSCIENCES INC. (ELIL ) LILLY & CO ELI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91WO-US07290.
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30-APR-1992.

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                                                                                                                                   Dovey HF, Johnstone EM, Little SP, McConlogue L, Seubert PA;
Sinha S;
                                                                                                                                                                                                                Human amyloidin protease - used for cleaving Met-Asp bond in amyloid-like substrate for identifying protease inhibitors
                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 60.0%; Score 6; DB 13; Length 10; Best Local Similarity 100.0%; Pred. No. 0.99; Matches 6; Conservative 0; Mismatches 0; Indels
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                                                                                                (ELIL ) LILLY & CO ELI.
                                                                                                                                                                                                                                                            Claim 1; Page 52; 62pp; English.
                                                    90US-0594122.
91US-0766351.
                          91WO-US07290.
                                                                                                                                                                                    WPI; 1992-167148/20.
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GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	Human APP derived Human amyloid prec	Human APP derived Human APP derived Kuman APP derived	Human APP derived N terminus of beta Human APP derived	Human APP derived Human APP A-beta p
SUMMARIES		AAB46213 AAB46215 AAB46212	AAB46216 AAU78518 AAB46211	AAB46217 AAB46202
80 80	15	222	53 53 53	22
å Query e Match Length DB	10	10	10 10 10	10
% Query Match	100.0	0.00	80.0 70.0 70.0	70.0
Score	10	J O B	7 7	7
Result No.	177	ມ 4 ເບ	9 7 8	9

Synthetic peptide Beta-amyloid pepti Human APP derived Human APP derived Beta amyloid pepti Beta-amyloid pepti Alpha secretase cl Doubly phosphoryla P.T.Aappa-B-alpha Human APP derived Beta-amyloid pepti Sphingolipid desat Beta-APP alpha-sec Human APP-Sw mutan Beta-amyloid pepti Epitope #1 used in Human amyloid beta Peptide derived fr Nucleotide-binding Immunogen for rais Immunogen for rais Humanised ATR-5 H Sphingolipid desat CDR region of anti Beta-secretase rel Laminin receptor-b Human amyloidin pr). ; clearing response; nootropic; seasesease. (T; c T; s administering amyloid .
7 23 AAO14421 8 19 AAW70865 10 22 AAB46210 10 22 AAB46210 5 23 ABB05182 6 19 AAW70868 6 23 AAW70868 10 21 AAY96712 10 22 AAB46219 10 22 AAB46219 4 19 AAW70870 5 22 AAB46219 6 22 AAB60636 5 22 AAB60636 6 22 AAB61099 6 22 AAB41099 6 22 AAB41099 7 21 AAB75340 8 20 AAW75340 9 22 AAU02741 9 23 AAB06519 10 13 AAR808279	lard; peptide; 10 AA. (first entry) ved immunogenic peptide #10. tt; APP; Abeta; brain; human; ediated phagocytosis; immunoger: sor protein; Alzheimer's dise; 2. 2. And F. Vasquez NJ, Yednock T ard F, Vasquez NJ, Yednock T treating a disease associated theimer's disease, comprises a cody
11111111111111111111111111111111111111	46214 stand 46214; APR-2001 (an APP deri: Loid deposi receptor me- loid precur 00072880-A2 DEC-2000. AMY-1999; UR-) NEURAL enk DB, Ba ; 2001-0321
	RESULT 1 AAB46214 ID AAB46214 ID AAB XX AAC AAB XX

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Conservative
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                                This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta; or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nociropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The methods are useful for prophylactic and therapeutic treatment of
                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beta amyloid is synthesised as part of a larger protein referred to as amyloid precursor protein (APP), which has a number of isoforms in humans, including APP65s and APP770. The amino terminal of beta amyloid is generated by cleavage of a peptide bond of APP which in APP695 lies between Met596 and Asp597. APP forms a complex with Go, a GTP-binding protein (or "G protein") in brain. Go is made of one alpha subunit and one Beta-agamma subunit. Two isoforms of Go, known as Gol (or GoA) and Go2 (or GoB) have been identified; they have slight AA differences in their alpha subunits. The cDNA sequence and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying cpds. useful for treating or preventing Alzheimer's disease. by determining whether it interferes with the association of the couplone portion of amyloid precursor protein to G polypeptide
                                                                                                                                                                                                                                                                 ;
0
                                                                                                                                                                                                                                        Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amyloid precursor protein; isoform APP 695; beta amyloid; Alzheimer's disease.
                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human amyloid precursor protein APP695 residues 597-606.
                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                     100.0%; Score 10; DB 22;
100.0%; Pred. No. 3.6e-05;
iive 0; Mismatches 0;
          Disclosure; Figure 19; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 33; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                             AAR58928 standard; peptide; 10 AA.
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                                                                                                                                                                                                                                                                 10; Conservative
                                                                                                                                                                                       Alzheimer's disease.
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                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                        1 AEFRHDSGYE 10
                                                                                                                                                                                                                                                                                                        10 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
deduced AA sequence of the alpha subunits in each of Gol and Go2 are shown in AAQ69002/R58914 and AAQ69004/R58924 respectively. The cytoplasmic App695 sequence His657-Lys676 (AAR58913) possesses a specific Go-activating function, and is necessary for complex formation of this APP with Go. AAR56928 is another peptide of AAP6695 which corresp. to AAR 597-606.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a disease associated with amyloid deposits, disease, comprises administering amyloid
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                                                                                                                                                                                                                                                                    Score 9; DB 15; Le
Pred. No. 0.00049;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human APP derived immunogenic peptide #9.
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100.0%; Pred. No.
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100.0%; Pre
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Best Local Similarity 100.
Matches 9; Conservative
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Gaps

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Indels

Mismatches

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26-MAY-2000; 2000WO-US14810.
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Matches 8; Conservative
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                                                                                   Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient. (a) an antibody that blinds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (Fc receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta; or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Alzheimer's disease.
                                                                                                                                                                    Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                               Yednock T;
                                                                                                                                                  Human APP derived immunogenic peptide #11.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Figure 19; 143pp; English.
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                                                                                  AAB46215 standard; peptide; 10 AA
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Best Local Similarity 100.
Matches 9; Conservative
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disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody that binds to Abeta, the antibody binds to an amyloid deposits of the brain of receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta; or (c) an agent that induces an immunogenic response against residues 1-3 of Abeta or (c) an agent at induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Alzheimer's disease.
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                                                                                    Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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Human APP derived immunogenic peptide #8.
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                                                                                                                        Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody ^{\rm -}
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                                                                                  Yednock T;
                                                                                                                                                                 Disclosure; Figure 19; 143pp; English.
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                                                                                 Vasquez NJ,
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                   26-MAY-2000; 2000WO-US14810.
                                        99US-0322289
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                                                            (NEUR-) NEURALAB LID.
                                                                                 Schenk DB, Bard F,
                                                                                                   WPI; 2001-032104/04.
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This invention relates to a novel method for modulating activity of beta-site amyloid precursor protein (APP)-cleaving enzyme (BACE)

beta-site amyloid precursor protein (APP)-cleaving enzyme (BACE)

consisted a soluble BACE which enhances the production of the enzyme results in the generation of a soluble BACE which enhances the production of the enzyme ratio of a soluble BACE which enhances the production of the enzyme ratio of an antisense nucleotide molecule capable of hybridishing with BACE mRNA, by using a ribozyme that targets and degrades BACE secretase. Inhibition of BACE secretase can be noting of the enzyme with BACE mRNA, by using a ribozyme that targets and degrades BACE secretase mRNA, with a peptide that can interfere with binding of the enzyme with BACE or using an antibody or antagonist that can innertion as an inhibitor of BACE secretase activation. The methods of the invention modulate the activity of BACE secretase/sheddase by preventing cleavage of BACE, which is useful for the treatment of a protein, especially Alzheimer's disease. The invention also comprises a method for identification of an agent that can alter the ability of BACE can be used for high throughput screening of candidate molecules. The invention also comprises a method for determining whether an individual is at risk of developing a neurodegenerative disorder characterised by the generation of Abeta protein by measuring the levels of BACE or forminal cleavage products in a sample or rissue where an increase consumption in the present sequence represents the N terminal of a beta amyloid protein of the
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Modulating activity of beta-site amyloid precursor protein-cleaving enzyme secretase/sheddase for treatment of neurodegenerative disorder characterised by generation of Abeta protein, by preventing cleavage of
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100.0%; Pred. No. 7.8e+05;
.ive 0; Mismatches 0;
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                                                                                                                                                                                                                         Disclosure; Page 28; 64pp; English.
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                                                                                                                    This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta; or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Mitheimer's disease.
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Pred. No. 0.09
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                                                                                             Disclosure; Figure 19; 143pp; English.
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100.0%; Pro
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antibody binds to an amyloid deposit and induces a clearing response (Foreceptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The methods are useful for prophylactic and therapeutic treatment of
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iive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-032104/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bard F,
                                                                                                                                                                                                                                                                                                                           Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                  10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 RHDSGYE 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200072880-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RHDSGYE 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HOMO sapiens.
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Gaps

.; 0

Indels

7.8e+05; hes 0;

Pred. No. 7.8 Mismatches

;

100.08;

Best Local Similarity 100. Matches 6; Conservative

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The invention relates to the field of the neurodegenerative disorder of Alzahamer's disease (AD). In particular, the invention provides a novel mutation (T7141) identified in the amyloid precursor protein (APP), and thich leads to a very aggressive form of AD. The mutation involves the 43rd coden of the amyloid Beta peptide (A-Beta) for involves the 43rd coden of the amyloid Beta peptide (A-Beta) for corresponding to the putative gamma 42-secretion elevating the A-Beta42 A-Beta40 and A-Beta40 and A-Beta42 secretion elevating the A-Beta42 A-Beta40 and A-Beta40 and A-Beta42 secretion elevating the A-Beta41 A-Beta40 and A-Beta40 and A-Beta42 benserored plaque pathology in brains of these patients is of the diffuse or pre-amyloid 'type composed primarily of N-truncated A-Beta42 benserored plaque cores, were composed primarily of N-truncated A-Beta42 for senile instance, in vessels as cerebral amyloid angiopathy (CAA) or senile instance, in vessels as cerebral amyloid angiopathy (CAA) or senile indicate that deposition of N-truncated A-Beta42 form. Together, these indicate that deposition of N-truncated A-Beta42 in one of the earliest amyloid deposition of N-truncated A-Beta42 in one of the earliest of inciting AD either through the well-setablished 'amyloid cascade' or by a yet unknown mechanism(s). This sequence represents a synthetic peptide of A-Beta42 by immunising mice with the synthetic peptide.

A-Beta42 by immunising mice with the synthetic peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neurodegenerative disorder; Alzheimer's disease; AD; T7141; APP; APP714; amyloid precursor protein; amyloid Beta peptide; A-Beta; A-Beta40; brain; A-Beta42; plaque pathology; pre-amyloid; cerebral amyloid angiopathy; Dense-cored plaque; CAA; senile plaque core; amyloid cascade; murine; mouse; DAEFRHD; moncolomal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel polynucleotide sequence encoding a mutant of amyloid precursor protein 70, useful for screening for a molecule capable of reducing the formation of beta amyloid 42 peptide \,
                                                                             Gaps
                                                                             ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          De Jonghe C, Kumar Singh S, Van Broeckhoven C;
                                                                     0; Indels
    Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic peptide of A-Beta residues 1-7 (DAEFRHD).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG.
60.0%; Score 6; DB 22; Le
100.0%; Pred. No. 7.8e+05;
1ve 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 21; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                          AA014421 standard; Peptide; 7 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUL-2001; 2001WO-EP07830.
    60.09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JUL-2000; 2000EP-0202362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                         Conservative
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Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200202769-A1.
                                                                                                                                           1 AEFRHD 6
                                                                                                                                                                                        111111
2 AEFRHD 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAY-2002
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA014421;
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                                                                                                                                                                                                                                                                                                                            RESULT 11
AAO14421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME OF THE PROPERTY OF THE PR
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DB 23;

Score 6;

60.08;

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The present sequence represents a peptide derived from beta-amyloid
precursor protein (beta-APP, see AAM70863). The peptide is a
precursor protein (beta-APP, see AAM70863). The peptide is a
beta-amyloid peptide and is used to produce a monoclonal antibody
designated antisemilin NJ. The specification describes a method for
prevention or inhibition of progression of Alzheimer's disease. The nethod comprises administering a composition comprising a recombinant bit of the N-terminus or the C-terminus of an amyloid-beta peptide, operably
inked to a promoter which is expressed in the central nervous system.
The recombinant antibody molecules prevent the accumulation of beta amyloid peptides in the extracellular space, interstitial fluid and the agraedation of such peptides into amyloid deposits in the brain. They also inhibit the progression of Alzheimer's disease by inhibiting the interaction of beta amyloid peptides mediating the interaction of beta amyloid peptides mediating consists in the complement activation and cytokine release involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Preventing or inhibiting progression of Alzheimer's Disease comprises use of recombinant DNA encoding an antibody specific for the N. or C-terminus of an amyloid-beta peptide
                                                                                                                                                                                                                                                   Beta-amyloid precursor protein; beta-APP; beta-amyloid peptide; antibody; amyloid deposit; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       60.0%; Score 6; DB 19; Length 8; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                 Beta-amyloid peptide to create a monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 46; 58pp; English.
                                                                                                                 AAW70865 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                          98WO-US06900.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          97US-0041850.
                                                                                                                                                                                 04-FEB-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the inflammatory process.
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nes 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MCIN/) MCINNIS P.A. (MIND-) MINDSET LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 AA;
1 AEFRHD 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 AEFRHD 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              111111
2 AEFRHD 7
                    W09844955-A1.
                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                          09-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                          15-OCT-1998.
                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                   AAW70865;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chain DG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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Matches
                                                                                   RESULT 12
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This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient. (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (Fc receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The methods are useful for prophylactic and therapeutic treatment of Alzheimer's disease.
Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beta amyloid peptide; beta-AP; beta amyloid precursor protein; A-beta, APP-770; amyloid aggregation; amyloidogqenic; Alzheimer's disease; mocorcopic; neuroprotective; immunosuppressive; antimicrobial; auditory; antidiabetic; antipyretic; dermatological; cardiovascular; nephrotropic; amyloid aggregation inhibitor; neurotoxatichy inhibitor; Down's syndrome; amyloidogenic disease; beta amyloid deposition; amyloidosis; hereditary cerebral haemorrhage; familial amyloid polyneuropathy.
                                                                                                                                                                                                                                                                                                                                                                          Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Beta amyloid peptide related peptide PPI-339 SEQ ID NO:36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22;
. 1.2;
                                                                                                                                                                                                                                                                                                  Bard F, Vasquez NJ, Yednock T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60.0%; Score 6; DB 2
100.0%; Pred. No. 1.2
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Figure 19; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABB05182 standard; peptide; 5 AA.
                                                                                                                                                                              26-MAY-2000; 2000WO-US14810.
                                                                                                                                                                                                                     99US-0322289
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                                                                                                                                                                                                                                                         (NEUR-) NEURALAB LTD.
                                                                                                                                                                                                                                                                                                                                      WPI; 2001-032104/04
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                                                                                               WO200072880-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 HDSGYE 10
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                                                             Homo sapiens.
                                                                                                                                                                                                                     28-MAY-1999;
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                                                                                                                                        07-DEC-2000.
                                                                                                                                                                                                                                                                                               Schenk DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABB05182;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient. (a) an antibody that blinds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Alzheimer's disease.
                                                                                                                                                                                                              Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Rc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yednock T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 6; DB 22
Pred. No. 1.2;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human APP derived immunogenic peptide #14.
                                                                                                                                                                              Human APP derived immunogenic peptide #6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Figure 19; 143pp; English.
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100.0%; Pre
                                                          AAB46210 standard; peptide; 10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vasquez NJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB46218 standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAY-2000; 2000WO-US14810.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0322289.
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                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NEUR-) NEURALAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-032104/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bard F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 especially Alzheim
specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200072880-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 AEFRHD 6
                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-APR-2001
                                                                                                                                        04-APR-2001
                                                                                                                                                                                                                                                                                                                                                                               07-DEC-2000.
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9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schenk DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S.
Matches 6,
                                                                                                 AAB46210;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB46218;
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                                    AAB46210
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Gaps

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Length 10; 0; Indels

20-NOV-2001

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The present invention describes an amyloid modulator compound (I) comprising an aggregation core domain and a modifying group attached to it. (I) has nootropic, neuroprotective, immunosuppressive, antimicrobial, antidiabbetic, antipyretic, dermatological, cardiovascular, nephrotropic, and auditory activities, and can be used as a natural amyloid aggregation inhibitor and a neurocoxicity inhibitor of natural bact amyloid aggregation inhibitor and an encrocoxicity inhibitor of natural amyloid aggregation inhibitor and an encrotrate of a medicament for the diagnosis or treatment of an amyloidogenic disease e.g. Alzheimer's disease and other clinical occurrences of beta amyloid deposition such as bown's syndrome individuals and in patients with hereditary cerebral haemorrhage with amyloidosis, and for treating a disorder associated with amyloidosis such as familial amyloid polymeuropathy. (I) reduces the amyloidosis to reduce the neurotoxic aggregates to cultured neuronal calls. (I) not only reduces the formation of neurotoxic aggregates but also have the ability to reduce the neurotoxicity of performed A-beta fibrils. The exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                        A, Benjamin H, Garnick MB, Gefter ML, Hundal A, Kasman L; Signer ER, Wakefield J, Reed MJ;
                                                                                                                                                                                                                                                                                                         Amyloid modulator compound useful for treatment of an amyloidogenic disease such as Alzheimer's disease comprises an aggregation core domain and a modifying group attached to it .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
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50.0%; Score 5; DB 23; Le
Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 5; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                  Example 11; Column 63; 54pp; English.
                                                                           95US-0404831.
95US-0475579.
95US-0548998.
                                        96US-0617267.
                                                                                                                                                                (PRAE-) PRAECIS PHARM INC.
                                                                                                                                                                                                                                                                        WPI; 2002-146668/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seguence 5 AA;
                                                                           14-MAR-1995;
07-JUN-1995;
27-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 HDSGY 9
                                     14-MAR-1996;
                                                                                                                                                                                                          Findels MA,
                                                                                                                                                                                                                                Musso G,
δ
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Search completed: April 21, 2003, 12:10:01 Job time: 28.3333 secs

1 HDSGY 5

Ωp

product of

Perfect score:

Sequence:

OM protein

Run on:

Scoring table:

Word size Searched:

Database :

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Synthetic oligopep
Doubly phosphoryla
p.1 kappa-B-alpha
Beta-APP alpha-sec
                                                                                                                                                            Beta-secretase rel
Human amyloidin pr
Beta-secretase sub
                                                                                                                     Human APP A-beta 1
Human amyloid beta
                                                           Human APP-Sw mutan
Human Asp-2 beta-s
                                                                                        Human beta-amyloid
Beta amyloid pepti
Beta-amyloid pepti
                                                                                                                                          Immunogen for rais
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                             Human APP derived
Human APP derived
Beta-APP alpha-sec
Human APP derived
Human APP derived
Beta-amyloid pepti
                                                                                                                                                                                                                                                                                                                                              Human APP-Sw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying cpds. useful for treating or preventing Alzheimer's disease - by determining whether it interferes with the association of the couplone portion of amyloid precursor protein to G polypeptide
                                                                                Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amyloid precursor protein; isoform APP 695; beta amyloid;
Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human amyloid precursor protein APP695 residues 597-606.
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                                                                                                                                                     AMI19507
ABB06519
ARR2465
AAR24260
AAR24260
AAR24261
AAR31756
AAR33756
AAR39766
AAR69709
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AAE10653
AAB46210
AAB46216
AAW70868
                             AAB46209
AAB46217
AAE10667
AAE06907
AAU06636
AAU0636
AAB05185
AAM70864
AAB46199
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AAB46199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93US-0019208
  15-APR-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GEHO ) GEN HOSPITAL CORP.
  WPI; 1994-294486/36.
  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nishimoto I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9419692-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-SEP-1994.
                                AAR58928;
                                                                                                                                                                                                                                                                                                                                                                                                          Human APP derived
Human APP derived
Human APP derived
Human APP derived
N terminus of beta
Human APP derived
Human APP A-beta
Synthetic peptide
Beta-amyloid pepti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                       April 21, 2003, 12:06:55; Search time 28.3333 Seconds (without alignments) 47.030 Million cell updates/sec
                                                                                                                                                                                                                                                                                                     1. (SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:*
2. (SIDSZ/gcgdata/geneseq-embl/AA1981.DAT:*
3. (SIDSZ/gcgdata/geneseq-embl/AA1981.DAT:*
4: (SIDSZ/gcgdata/geneseq-embl/AA1981.DAT:*
5: (SIDSZ/gcgdata/geneseq-embl/AA1983.DAT:*
6: (SIDSZ/gcgdata/geneseq-embl/AA1985.DAT:*
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10: (SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1990.DAT:*
11: (SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1990.DAT:*
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20: (SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1990.DAT:*
21: (SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1990.DAT:*
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23: (SIDSZ/gcgdata/geneseq/geneseqp-embl/AA2000.DAT:*
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             GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                     908470 seqs, 133250620 residues
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AAB46213
AAB46212
AAB46214
AAU78518
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AAB46202
AAO14421
AAW70865
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                                                    - protein search, using sw model
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Gapop 60.0 , Gapext 60.0
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Match
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Score

Result No.

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Gaps

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Indels

Length 10;

100.0%; Score 10; DB 22; 100.0%; Pred. No. 3.5e-05; ive 0; Mismatches 0;

Conservative

10 AA;

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that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The methods are useful for prophylactic and therapeutic treatment of Alzheimer's disease.
                                                                                                                        Query Match
Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                   1 DAEFRHDSGY 10
                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                    Beta amyloid is synthesised as part of a larger protein referred to as amyloid precursor protein (APP), which has a number of isoforms in humans, including APP695 and APP770. The amino terminal of beta amyloid is generated by cleavage of a peptide bond of APP which in APP695 lies between Met596 and Asp597. APP forms a complex with Go, a GTP-binding protein (or "G protein") in brain. Go is made of one alpha subunit and one Beta-gamma subunit. Two isoforms of Go, known as Gol (or GoA) and Go2 (or GoB) have been identified; they have slight AA differences in their alpha subunits in each of Gol and Go2 are shown in AAQ69002/R58914 and AAQ69004/R58924 respectively. The cybolasmic App695 sequence His657-Lys676 (AAR58913) possesses a cycoplasmic App695 sequence His657-Lys676 (AAR58913) possesses a formation of this APP with Go. AAR58928 is another peptide of APP695 which corresp. to AAR 597-606.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta; or (c) an agent
                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody
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                                                                                                                                                                                                                                                                                              100.0%; Score 10; DB 15;
100.0%; Pred. No. 3.5e-05;
1ve 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                            0; Mismatches
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             Disclosure; Page 33; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB46213 standard; peptide; 10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vasquez NJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAY-2000; 2000WO-US14810.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0322289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                           10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                      1 DAEFRHDSGY 10
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                                                                                                                                                                                                                                                                     10 AA;
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                                                                                                                                                                                                                                                                       Seguence
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This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (For receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at lasat residues 1-5 of Abeta; or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment methods are useful for prophylactic and therapeutic treatment of Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                 Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid
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100.0%; Pred. No. 0.0
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yednock
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                                                                                                                                                                           AAB46212 standard; peptide; 10 AA.
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Best Local Similarity 100.
Matches 9; Conservative
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specific antibody
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This invention relates to a novel method for modulating activity of beta site amyloid precursor protein (APP)-cleaving enzyme (BACE) beta site amyloid precursor protein (APP)-cleaving enzyme (BACE) secretase/sheddase. Cleavage of BACE by this enzyme results in the generation of a soluble BACE which enhances the production of the amyloidogenic peptide Abeta which has been shown to be involved in the actionary of Alzheimer's disease. Inhibition of BACE secretase can be actionary of Alzheimer's disease. Inhibition of BACE secretase can be actionary of Alzheimer's sing a ribosyme that targets and of hybridising with BACE mRNA, by using a ribosyme that targets and binding of the enzyme with BACE mRNA, by using a ribosyme that targets and function as an inhibitor of BACE secretase activation. The methods of the invention modulate the activity of BACE secretase/sheddase by preventing cleavage of BACE, which is useful for the treatment of a neurodegenerative disorder characterised by the generation of Abcta protein, especially Alzheimer's disease. The invention also comprises a method for identification of an agent that can alter the ability of BACE can be used for high throughout screening of candidate molecules. The invention also comprises a method for determining whether an individual is at risk of developing a neurodegenerative disorder characterised by the generation of Abcta protein by measuring the levels of BACE charminal cleavage products in a sample or tissue where an increase comparage products in a beta amyloid protein of the protein of the protein of the aperson at risk. The present sequence represents the N terminal of a beta amyloid protein of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modulating activity of beta-site amyloid precursor protein-cleaving enzyme secretase/sheddase for treatment of neurodegenerative disorder characterised by generation of Abeta protein, by preventing cleavage of
                                                                                           Alzheimer's disease; beta amyloid precursor protein; mouse; BACE; beta-site APP cleaving enzyme; nootropic; neuroprotective; beta-site amyloid precursor protein (APP)-cleaving enzyme; APP; BACE secretase/sheddase; neurodegenerative disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       80.0%; Score 8; DB 23; L6 100.0%; Pred. No. 7.8e+05; iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                         01-AUG-2001; 2001WO-CA01118
                                                                                                                                                                                                                                                                                                                                                                                                            01-AUG-2000; 2000CA-2313828
                                             N terminus of beta amyloid.
18-JUN-2002 (first entry)
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es 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 DAEFRHDS
                                                                                                                                                                                                                                                                                                              07-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seidah NG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .nvention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                     Mus sp.
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Matches
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AAB46211
ID AAB46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Nizheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                    Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; for receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.0%; Score 9; DB 22; Length 10; 100.0%; Pred. No. 0.00048; rive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vasquez NJ, Yednock T;
                                                                                                                                                                                                                                                                                                                                         Human APP derived immunogenic peptide #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Figure 19; 143pp; English.
                                                                                                                                                                                            AAB46214 standard; peptide; 10 AA
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                                                                                                                                                                                                                                                                                        (first entry)
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Best Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 AEFRHDSGY 10
                                                     10 AA;
                        1 DAEFRHDSG 9
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                                                                                                                                                                                                                                          AAB46214;
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Gaps

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0; Indels Length 8;

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WO200072880-A2.

07 - DEC - 2000

Homo sapiens.

04-APR-2001

28-MAY-1999;

Schenk DB,

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This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                                                                                                                                                                                         Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 8; DB 22; Length 10;
Pred. No. 0.0066;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human APP A-beta protein N-terminal fragment.
                                                                                                                                                                                                              Yednock T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yednock T;
                                                                                                                                                                                                                                                                                                                                                                             Disclosure, Figure 19; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   80.0%; Scc.
100.0%; Pred
0; F
                                                                                                                                                                                                              Vasquez NJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB46202 standard; peptide; 7 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-MAY-2000; 2000WO-US14810.
                                                                                    26-MAY-2000; 2000WO-US14810.
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Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                    (NEUR-) NEURALAB LID
                                                                                                                                                                                                           Schenk DB, Bard F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NEUR-) NEURALAB LID.
                                                                                                                                                                                                                                                      WPI; 2001-032104/04.
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WO200072880-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200072880-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-MAY-1999;
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                                            07-DEC-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB46202;
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AAB46202
    δλ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis against it (b) a polypeptide containing an N-terminal segment of at least residues 1.5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1.3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The Abetament's disease.
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                                                                                                                       Amyloid deposit; App; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzabimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yednock T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human APP derived immunogenic peptide #11.
                                                                               Human APP derived immunogenic peptide #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Figure 19; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB46215 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vasquez NJ,
                                                                                                                                                                                                                                                                                                                              26-MAY-2000; 2000WO-US14810. ·
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                                       (first entry)
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nes 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                   (NEUR-) NEURALAB LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bard F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-032104/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 AA;
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1 DAEFRHDS 8

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Sequence

Query Match Best Loc Matches Homo saplens

04-APR-2001

AAB46215;

AAB46215 RESULT

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Gaps

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Alzheimer's disease (AD). In particular, the invention provides a novel mutation (T7141) identified in the amyloid precursor protein (APP).

APP714, which leads to a very aggressive form of AD. The mutation involves the 47d codon of the amyloid Beta petide (A-Beta)

Corresponding to the putative gamma 42-secretase cleavage site. The cover mutation alters both A-Beta40 and A-Beta42 secretion elevating the A-Beta42/A-Beta40 ratio by 10-fold in vitro. Furthermore, the main amyloid plaque pathology in brains of these patients is of the diffuse of pre-amyloid type composed primarily of N-truncated A-Beta42. Dense-cored plaques although not absent, were significantly reduced. Also, the usual sites in brain where A-Beta40 is predominantly deposited, for instance, in vessels as carebral amyloid anglopathy (CAA) or senile classes were composed entirely of A-Beta42 form. Together, these indicate that deposition of N-truncated A-Beta42 in one of the earliest of manyloid deposited in the brain, the diffuse plaques, is fully competent of inciting AD either through the well-established 'amyloid cascade' or by a yet unknown mechanism(s). This sequence represents a synthetic peptide of A-Beta4, residues 1-7 (DAEFRHD). This sequence was used for the arising a monoclonal antibody specific for the N-terminus of A-Beta40 and A-Beta42 by immunising mice with the synthetic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a peptide derived from beta-amyloid precursor protein (beta-APP, see AAW70863). The peptide is a beta-amyloid peptide and is used to produce a monoclonal antibody designated antisenilin NI/7. The specification describes a method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Preventing or inhibiting progression of Alzheimer's Disease - comprises use of recombinant DNA encoding an antibody specific for the N- or C-terminus of an amyloid-beta peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Beta-amyloid precursor protein; beta-APP; beta-amyloid peptide; antibody; amyloid deposit; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 23; Length 7; 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beta-amyloid peptide to create a monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          70.0%; Score 7; DB 2
100.0%; Pred. No. 7.8
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW70865 standard; peptide; 8 AA.
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Best Local Similarity 100..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                         This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The methods are useful for prophylactic and therapeutic treatment of Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neurodegenerative disorder; Alzheimer's disease; AD; T7141; APP; APP714; amyloid precursor protein; amyloid Beta peptide; A.Beta; A.Beta40; brain; A.Beta42; plaque pathology; pre-amyloid; cerebral amyloid angiopathy; bense-cored plaque; CAA; senile plaque core; amyloid cascade; murine; mouse; DAEFRHD; monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel polynucleotide sequence encoding a mutant of amyloid precursor protein 70, useful for screening for a molecule capable of reducing the formation of beta amyloid 42 peptide \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to the field of the neurodegenerative disorder of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                     Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 7; DB 22; Length 7;
Pred. No. 7.8e+05;
0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic peptide of A-Beta residues 1-7 (DAEFRHD).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kumar Singh S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 21; 42pp; English.
                                                                                                                                                    Page 119; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  70.05; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA014421 standard; Peptide; 7 AA.
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Best Local Similarity 100...
7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-154742/20.
          WPI; 2001-032104/04
                                                                                                       specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200202769-A1.
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1 DAEFRHD 7
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Synthetic.
                                                                                                                                                    Claim 59;
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Gaps

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us-09-580-018-9.oli.rag

Alzheimer's disease.

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prevention or inhibition of progression of Alzheimer's disease. The method comprises administering a composition comprising a recombinant bwa molecule containing a gene encoding a recombinant antibody end-specific for the N-terminus or the C-terminus of an amyloid-beta peptide, operably linked to a promoter which is expressed in the central nervous system. The recombinant antibody molecules prevent the accumulation of beta-amyloid peptides in the extracellular space, interstitial fluid and cerebrospinal fluid and the aggregation of such peptides into amyloid deposits in the brain. They also inhibit the progression of Alzheimer's disease by inhibiting the interaction of beta-amyloid peptides mediating Alzheimer's disease induced neurotoxicity and inhibiting the Alzheimer's disease induced complement activation and cytokine release involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (For receptor mediated phagocytosis) against it (b) a polyeptide containing an N-terminal segment of at least residues 1-5 of Abeta: or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The products of the invention have nootropic and neuroprotective activity. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The methods are useful for prophylactic and therapeutic treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody
                                                                                                                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                                                                                                                                                                                            \prime Match 70.0%; Score 7; DB 19; Length 8; Local Similarity 100.0%; Pred. No. 7.8e+05; hes 7; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human APP derived immunogenic peptide #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Figure 19; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB46210 standard; peptide; 10 AA.
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                                                                                                                                                                                                                                                           the inflammatory process.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 1 DAEFRHD 7
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Best Local S:
Matches 7,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amyloid deposit; APP, Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
                                                                                                                    Gaps
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No. 0.091;
0; Indels
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                                                    Score 7; DB 2; Pred. No. 0.0 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human APP derived immunogenic peptide #12.
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                                                                                                                                                                                                                                                                                                                                                                                                       AAB46216 standard; peptide; 10 AA.
                                                    70.0%;
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                                                            Query Match 70.0
Best Local Similarity 100.
Matches 7; Conservative
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Best Local Similarity
Matches 7; Conserv
10 AA;
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Sequence
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(NEUR-) NEURALAB LID.
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|5 DAEFRH 10
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                                                                                                                                                      WO200072880-A2.
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                                                                                                                       Homo sapiens.
                                                                                                                                                                                    07-DEC-2000.
                                                                                                                                                                                                                                                                                                              Schenk DB,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a peptide derived from beta-amyloid precursor protein (beta-APP, see AAW/0863). The peptide is a precursor protein (beta-APP, see AAW/0863). The peptide is a percursor protein (beta-APP, see AAW/0863). The peptide is a pecufication describes a method for prevention or inhibition of progression of Alzheimer's disease. The method comprises administering a composition comprising a recombinant DNA molecule containing a gene encoding a recombinant antibody end-specific for the N-terminus or the creminus of an amyloid-beta peptide, operably linked to a promoter which is expressed in the central nervous system. The recombinant antibody molecules prevent the accumulation of beta-amyloid peptides in the extracellular space, interstitial fluid and cerebrospinal fluid and the agreedation of such peptides into amyloid deposits in the brain. They also inhibit the progression of Alzheimer's disease by inhibiting the interaction of beta-amyloid peptides mediating Alzheimer's disease induced complement activation and cytokine release involved in the inflammatory
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                                                                                                                                                                  Beta-amyloid precursor protein; beta-APP; beta-amyloid peptide; antibody; amyloid deposit; Alzheimer's disease.
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                                                                                                                                        Beta-amyloid peptide to create a monoclonal antibody.
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                                          AAW70868 standard; peptide; 6 AA
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                                                                                                                                                                                                                                                                                                                                                            97US-0041850
                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                          (MCIN/) MCINNIS P A. (MIND-) MINDSET LTD.
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                                                                                                                                                                                                                                    Homo sapiens.
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                                                                           AAW70868;
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AAB46209
ID AAB46
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AC AAB46
               RESULT 13
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                                                                     Amyloid deposit; APP; Abeta; brain; human; clearing response; nootropic; Fc receptor mediated phagocytosis; immunogenic response; neuroprotective; amyloid precursor protein; Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bard F, Vasquez NJ, Yednock T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human APP derived immunogenic peptide #13.
Human APP derived immunogenic peptide #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Figure 19; 143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-MAY-2000; 2000WO-US14810.
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Matches 6; Conservative
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This invention describes a novel method of preventing or treating a disease associated with amyloid deposits of amyloid precursor protein (APP) Abeta fragments in the brain of a patient, which comprises administering to the patient: (a) an antibody that binds to Abeta, the antibody binds to an amyloid deposit and induces a clearing response (FC receptor mediated phagocytosis) against it (b) a polypeptide containing an N-terminal segment of at least residues 1-5 of Abeta, or (c) an agent that induces an immunogenic response against residues 1-3 to 7-11 of Abeta. The method is also useful for monitoring a course of treatment being administered to a patient e.g. active and passive immunization. The methods are useful for prophylactic and therapeutic treatment of Alzheimer's disease.
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                                                                                                                                                                                                             Preventing or treating a disease associated with amyloid deposits, especially Alzheimer's disease, comprises administering amyloid specific antibody
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 60.0%; Score 6; DB 22; Length 10; Best Local Similarity 100.0%; Pred. No. 1.2; Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                           Schenk DB, Bard F, Vasquez NJ, Yednock T;
                                                                                                                                                                                                                                                                                          Disclosure, Figure 19; 143pp; English.
                                26-MAY-2000; 2000WO-US14810.
                                                                    28-MAY-1999; 99US-0322289
                                                                                                      (NEUR-) NEURALAB LTD.
                                                                                                                                                                               WPI; 2001-032104/04.
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